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Fig.1A

SEQ:	ID	
NO1	50	
208	HCV-11aATGAGCACGAATCCTAAA	ACCTCAAAAAAAAAACAAACGTAACACCAACCG
209	HCV-J1bA	
210		GC
1		·GC
5		GC
9		GCA-A
11		GCC
211	HC-J62aA	GCA-A
212		A-AA
213		GCA-AT
214		GCA-AT
17		GCTA-AT
13		
21		GCA-AT
21	DNESZII A	0 C A A
215	NZI.13aACT	GCA-AT
216		G-CCA-AACT
217		-CGCA-AT
218		CGCA-AT
219		CGCA-AGT
220		G-CCCA-AACC
220	MEIZSSITTERITERITE	
221	744a	GC
222		GC
223		GC
224		GC
225		-TGC
27		GC
21	BNL/4K	G
226	DP0EE2	GCA-A
220	везоза	GCA-A
227	ΨΚ263λCTλ	CGCA-A
221	MKZ0aACIA	CGCA-A
228	EP172	CGCA-ATT
220	rki/aAciA	CGCA-A11
43	VN48aΔCTλC	GCA-AT
45		GCA
	111200 ACI	5C h
47	VN129aACTA	CGCA-AA
- '	THE ACT -A	C C - A-AA
49	NE9810aACT	-AGN
	HOTOLOG MCI	C AAN



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Fig.1B

SEQI		
NO51		
	CV-11aTCGCCCACAGGACGTCAAGTTCCCGG	
209H	CV-J1bCT	CT
	[C-G91cCTT	
	11dCTK-GSNNNNNNN	
	21dC	
9CAM	10781eC	CTC
11FR2	21fC	GGG
211H	[C-J62aATT	CC
212H	C-J82bC	
21358	832cC	CTC
	E922dC	
	42f	
	L32eC	
	L52hC	
		. 1 6
215	NZL13a	
216	HCV-TR3bAT	
217	NE483c	
218	NE2743dT	
219	NE1453eGAT	
220	NE1253fC	
220	NB12551C	C1G
221	Z44aCCATA	ТС
222	Z14bCATTGA	
223	GB3584cCCATT	
223		
225	DK134dCATT	
-	GB8094eCCATT	
27	BNL74kCCATT	TCC
226	DEOCC-	G
226	BE955a	CT
227	HK26aAC	•
227	HK26aAC	·
	DD 1 7	
228	FR17aTAT	CC
43	VN48aC	
45	VN138b	
1_		
47	VN129aATT	C
49	NE9810aCGT	· A C



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Fig.1C

SEQI			
NO10			000000000000000000000000000000000000000
	HCV-11aTTTACTTGTTGCCGCG		
	HCV-J1bC		
210	HC-G91c		
	L11dC		
	L21dC		
	M10781e-CGC-A		
11FR	21f	CG	G
211	HC-J62a-A		AG
212			
213			
214			
17	FR42f		
13	BNL32e		
21	BNL52h-A		
41	BNL52II-A		
215		AC	
216		TAC	
217	NE483c-AG		TAC-T
218	NE2743d-CAC	A	AGTTC-T
219	NE1453e-A	AC	ATC-T
220	NE1253f-AG-A	AC	AGT-C-T
221	Z44a	CG	TC
222	Z14bC		AG-TC-G
223	GB3584c		TG
224	DK134d		TG
225		G	
27	BNL74k		
2,	BALLIAN	6 6	10 0
226	BE955a	GA	TC-G
227	HK26a	CC-G	
228	FR1/a	C-T	
43	. VN48a-CC		
45	VN138b		G
47	VN129a-CA	AC-T	G
49	NE9810aGC-AA		TAGT-C-C



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Fig.1D

SEQID NO151200 208HCV-11aAAGACTTCCGAGCGGTCGCAACCTCGAGGTAGACGTCAGCCTATCCCCAA 209HCV-J1b
211HC-J62aGCGATAGCCT 212HC-J82bACGGTACC
215NZL13aATAAGCACA
221Z44a
226BE955aGA
228FR17aCACGACGCCAA
43VN48aT-AC-GCAGC-AAA 45VN138bAT-AC-GCA-GC-AAG
47VN129aGA
49NE9810aG



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Fig.1E

SEQID	
NO201250	
208HCV-11aGGCTCGTCGGCCCGAGGGCAGGACCTGGGCTCAGCCCGGGTACCC	TTGGC
209HCV-J1bCTT	
210HC-G91cCCAATG	
1BNL11dTTTT	
5BNL21dA-TTNNAC-TC-	
9CAM10781eAGCAT	
11FR21fA	· - -
211HC-J62aAGCTACTAATGAA-AAAA	C
212HC-J82bA-AGCTACCA-TGAAAAT	-
213S832cA-AGCAACTA-TGAAGAA	
214NE922dA-AGCACTA-TGAA-AAA	
17FR42fA-AGCGACTA-TGA-GTAA	
13BNL32eA-AGN-NGACTTGA-GTAATC-	
21BNL52hA-AGCTACTAATGA-GTAA	
ZIBNL5ZIIA-AGCIACIAAIGA-GIAA	
215117112	
215NZL13aGAGACT	
216HCV-TR3bCTCGCT	
217NE483cGTGGACTG	
218NE2743dAAGCTT	
219NE1453eAC-C-AGGAACTGT	
220NE1253fACAAGCTCT	
221Z44aGC-AA	
222Z14bGCTT	
223GB3584cAAT-TAT	
224DK134dGC-AA-TTTTT	
225GB8094eGCATATGT	
27BNL74kGATATAAATA-	
226BE955aGC-AACCTGA	
227HK26aGC-ACACA	
228FR17aTAC-AGACAC-T-GGAC	
43VN48aA-TGC-AC-AAACC-TC	
45VN138bTGAC-AAACC-TAAC-	
47VN129aTGC-A-AA-C-AC-A	
49NE9810aGCAATT	



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Fig.1F

NO251300 208HCV-11aCCCTCTATGGCAATGAGGGCTGCGGGTGGGCGGGATGGCTCCTGTCTCCC 209HCV-J1b
209HCV-J1b
210HC-G91c
1BNL11dC 5BNL21dC 11FR21fCTCAC 211HC-J62aACGACTCA
5BNL21dC 11FR21fCTCAC 211HC-J62aACGACTCA
11FR21fCTCA
211HC-J62aACGACTCAC 212HC-J82bGCACTC
211HC-J62aACGACTCAC 212HC-J82bGCACTC
212HC-J82bGCACTCTT
212HC-J82bGCACTCTT
17FR42fGCGCCTCAG
13BNL32eGGCTCA
21BNL52hGCTTTATCT
215NZL13aCA
216HCV-TR3bCGATTATC
217NE483cCT
218NE2743d-TTTATC
219NE1453eTCT
220NE1253fGTA
221Z44aT
222Z14bTCTAG
223GB3584c-TTCTTAT
224DK134dTCA
225GB8094eT-CTAGCT
27BNL74k-TTCT
2'26BE955aTC-CCTAGGCCT
227HK26a-TTACTAT
228FR17a
43VN48a-TTATTACC
45VN138b-TTGTTCAG
in the second se
47VN129aTGC
1, 1,1,1,2,0
49NE9810aACG
49NF98102>



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Fig.1G

SEQID	
NO301350	
208HCV-11aCGTGGCTCTCGGCCTAGCTGGGGCCCCACAGACCCCCGGCGTAGGTCGCG	,
209HCV-J1bTT	
210HC-G91cCTTT-T	
1BNL11dC	
5BNL21dC	
11FR21fCAAATAAAA	
211HC-J62aATCTCTCTATAAC	
212HC-J82bCGTCTCAAA	
213S832cCTCTCACAAA	
214NE922dAGCGTCAATACA	
17FR42fGAACAACAC	
13BNL32eA	
21BNL52hA	
215NZL13aCCTATCA-ATGC	
216HCV-TR3bTCTA-ATA-C-	-
217NE483cCTGA-ATA-AAC	
218NE2743dCATCTATCTATCT	
219NE1453eCAGTACAAC	
220NE1253fA-AATA-ATA-A	
221Z44aCATCTA-ATTGA	
222Z14bCTCAGTCTATT	
223GB3584cAGTCTA-ATTAC	
224DK134dGTCTG-ATTG-C	
225GB8094eCGGTCTT-ATTGC	
27BNL74kCT	
226BE955aAATATAAA	
227HK26aCC-ACATATC-AC	
220 FP 17 2 C C M 2M 2C 2 C	
228FR17aCGATACAC	
43VN48aCCA-ATA-ACGC	
45VN138b-NCCATT-ATN-GC	
47VN129a	
4.4M123dCGGMMM1M-GC	
49NE9810aC	
47NB70104 C	



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Fig.1H

SEGID	
NO351400	
208HCV-11aCAATTTGGGTAAGGTCATCGAT	'ACCCTTACGTGCGGCTTCGCCGACCTCA
209HCV-J1bT	
210HC-G91c	CT
11FR21f	ATT
211HC-J62aCG	
212HC-J82bC-GA	ATTT
213S832cC	A
214NE922dC	TT
17FR42fC	CTT-S
15BNL32eN-NT	
215NZL13aAA	AA
216HCV-TR3bCT	ATA
217NE483c	AG
218NE2743dCCA	
219NE1453eC	TCAGT
220NE1253fC	CTAT
221Z44aC	G
222Z14bTCA	GT
223GB3584cC	ACT
224DK134dC	AT
225GB8094eCC	
226BE955aT	AAT
227HK26aG	ATGT
228FR17aC	A-NNC-A
43VN48aCAC	T
45VN138bCC	TNS
47VN129aCC	CT



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Fig.11

SEQID NO401450 208HCV-11aTGGGGTACATACCGCTCGTCGGCGCCCCTCTTGGAGGCGCTGCCAGGGCC 209HCV-J1bT
211HC-J62aC-TGAG-C-C-CTCAT 212HC-J82bCGTGTGGTCAT 213S832cCGTT-CGCTA 214NE922dC-TGAGT-T-TCAT 17FR42fTGG-G-CTA 15BNL32eN-CG-T,GG-G-CG-TN
215NZL13a
221Z44aACAGCG-GGTCT 222Z14bAT
226BE955aTCAGCAGTCAT 227HK26aTCGGGT-GCTCGGCTG
228FR17aCTGC-AA-GGGCTGGCT
43VN48aTCTGATGW-GTCGGN 45VN138b-A-AT
47VN129aACTGTCTGGCAA



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Fig.1J

SEQID
NO451500
208HCV-11aCTGGCGCATGGCGTCCGGGTTCTGGAAGACGGCGTGAACTATGCAACAGG
209HCV-J1bAT
210HC-G91cATTA-ACTC
3BNL11d
7BNL21d
11FR21fN-ATCNGTNNNNNNNNNNNNNN
211HC-J62aCGA-ACGTT-T
212HC-J82bACTTACGA-ATC
213S832cCCGGA
214NE922dCGA-AGA-A
15BNL32eCNGG
17FR42fC
19BNL42gGAT 23BNL52hGA-AC
25BNL62iGA-A
25BNL621GA-A
215NZL13aCGACCTGA-AT-TC
216HCV-TR3bCTTGACAT-GGA
217NE483cCGACT
218NE2743dCATGA-A-CCTGAA-AT-TC
219NE1453eCACGAACCGAA-AT-T
220NE1253fAATGACTGAA-AT
221Z44a
222Z14bCACCGGAA-TC
223GB3584cACTTAC-GGA-CTG
224DK134dACGGCT
225GB8094eACTTAC-GGA-CC
29BNL74kGA-CT-T
31BNL84kGA-CT
33BNL94kGA-TT
35BNL104kGA-CT
37BNL114kGA-TT
39BNL1241GA-CT
226BE955aCACTGACTGGA
220bb955aCAC1GAC1GA
227HK26aCAGACAA-CGGA-CT
228FR17aTACAA-CGGCTC
43VN48aT
+3 VN + 0 a I G AN N CA - C G N A I C N
47VN129aNATACCA-CGGA-A
51NE9810aAA-TT-TC



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Fig.1K

SEQID		
NO501550		
208HCV-11aGAACCTTCCTGGT	TGCTCTTTCTCTA	TCTTCCTTCTGGCCCTGCTCTCTT
209HCV-J1bTGC		CT-ATTG
		T-GCTTAC-
3BNL11dT-GC		
7BNL21dTT-G		
11FR21fNN		C1N1-A
		GC-
		TT-GTTGA-
213S832cTT-GC		
214NE922dT-GC		
15BNL32eC		
17FR42fT-GC	CT	T-GTCT-G
19BNL42gTG		T-GTTG
23BNL52hTGC		
25BNL62iG		
215NZL13aT-GC	СТ	TTT
		TCCTCTC-
217NE483cTT-A		
218NE2/43Q11-AC		T-GTTT
219NE1453e	T	GTTGA-
220NE1253fTT-GC	CT	A-
221Z44aTC		
222Z14b		
223GB3584cTC		T-CTATTG-
224DK134dTC		G-
225GB8094eTCC	CT	G-
29BNL74kCC	CT	G-
31BNL84kC	T	CTACG-
33BNL94kTC	CT	G-
35BNL104kTAC		
37BNL114kYCC		
39BNL1241CC		
39BNL1241CC		A-CATG-
		A-CATG-
39BNL1241CC 226BE955aTT-AC		A-CATG-
39BNL1241CC		A-CATG-
39BNL1241CC 226BE955aTT-AC 227HK26aTCC		A-CATG- TATTTG- TAA
39BNL1241CC 226BE955aTT-AC		A-CATG- TATTTG- TAA
39BNL1241CC 226BE955aTT-AC 227HK26aTCC		A-CATG- TATTTG- TAA
39BNL1241CC 226BE955aTT-AC 227HK26aTCC	CT	A-CA-TGTATTTGTAAAG-
39BNL1241CC 226BE955aTT-AC 227HK26aTCC 228FR17aT	CT	A-CA-TGTATTTGTAAAG-
39BNL1241CC 226BE955aTT-AC 227HK26aTCC 228FR17aT	CT	A-CA-TGTATTTGTA-AGCT-AAT-AG-
39BNL1241CC 226BE955aTT-AC 227HK26aTCC 228FR17aT 43VN48aTCNN	CT	A-CA-TGTATTTGTA-AGCT-AAT-AG-
39BNL1241CC 226BE955aTT-AC 227HK26aTCC 228FR17aT 43VN48aTCNN	CT	A-CA-TGTATTTGTAAAGCT-AAT-AGNCTA-TG-



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Fig.1L

SEQID
NO551600
208HCV-11aGCTTGACTGTGCCCGCTTCGGCCTACCAAGTGCGCAACTCCACGGGGCTT
209HCV-J1b-TCA-CACTG-GGTGT-CA-A
210HC-G91cCACTGT-GGTTG-G
3BNL11dG-TAA-KA-CTCG-GG-AT-CG-G
7BNL21dG-TAAA-CTC-TG-GG-AT-CG-A
11FR21fC-CACTTG-GAG-A-AC-ATGGC
11FR211C-CAC11G-GAG-A-AC-A1GGC
211HC-J62aA-CCACCG-TCCTGC-GAAGATGTACCGGC
212HC-J82bG-CAA-TGTAGTGGCA-GATT-GTTCTAGC
213S832cA-CTA-TCGTGG-GCAAGGAGGC-ACTCC
214NE922d-TA-CG-TCC-GTGGCAAGAGCA-CTC-
15BNL32e-TG-CCT-TCT-N-GTTG-GCAAATAGTCA-GCC
17FR42f-TA-CCTGTATAGTAAGAAGCCACT-C
19BNL42g-TG-CCT-TCTGTGGTAAGAGTACCA-G
23BNL52h-TC-CGGCTGTGGCAAGAGCCACTC-
25BNL62iA-CCG-TCTGTGTGCGCGGTTTC-
215NZL13aA-T-CATAAG-CAGTCTAG-GTGGTA-GT-TCC
216HCV-TR3bTGCGT-GTAG-GTACACGA-GT-TCA
217NE483cGTCTGTTAG-A-GGCT-G-GTACGTGTAT-CCC
218NE2743dGTCTGTTG-A-GGATTGTACGTGTGT-TCC
219NE1453eCT-TGCTAGTC-GG-TGG-GTG-AT-CTC
220NE1253fGT-TCCAGGGCTAG-GTACA-GA-GT-CCA
220NE12331G1-1CCAGGGCIAG-G1ACA-GA-G1-CCA
221Z44aC-CTAGTG-GCTACGTG-TTCA-C
222Z14bCAACAAATGTGCTACGTG-TTCG-C
223GB3584cCTA-CGT-A-CTATTGTCA-C
224DK134dCTG-C
225GB8094eC-CTGG-GTTA-CTATTG-TTCG
29BNL74kCCA
31BNL84kCTATTA-CTACA-TCA-C
33BNL94kCCATTA-CTAC-AA-TCA-C
35BNL104k-TCCA-C
37BNL114kCCA
39BNL1241CCGCTC-GTTATGTGT-TCA
226BE955a-TCCTGCTAGTT-CCTACATGT-TA
227HK26aC-CAACATCTTACCTACGGTA
228FR17aC-CACAACAAATTCAAGGT-TA-C
43VN48aC-TAACAACCGGCGTTATACAAGT-TCG
47VN129aC-CCACTCCACTAA-CTATGCTAAGT-TG
51NE9810aCT-ACAA-AG-C-GGCTGG-GTACTTGT-CAC



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Fig.1M

SEQID NO601650 208HCV-11aTACCACGTCACCAATGATTGCCCTAACTCGAGTATTGTGTACGAGGCGGC 209HCV-J1bTGCCT-CAAAA 210HC-G91cTCCTT-CAAA 3BNL11d-T-TCCT-T-CCCA-CTAT-AG 11FR21fTTCTT-CGGCCCATAAA
211HC-J62aATGGCCA-CTGATCACC-GGC-ACTCCA
212HC-J82bTCTT-AAACCCACC-GGCCTCA-
213S832cATGCCGCT-CT-CT-GGCCTT-A
214NE922dATGACAGAGTCCC-GGCCTCAG
15BNL32eTATG-CACCT-CAACCCA-GGC-ATTN
17FR42fATG-CGTCTG-CTGACCCC-GGCCTCAG
19BNL42qATG-CACTT-CAACCCA-C-GGC-AAT-CA
23BNL52hTATGGT-AAGCCC-GGCCTTAA
25BNL62iATGGT-GAGCCCT-GGCCTC-A
10 1 10
215NZL13aGT-C-TCCTT-CTAGCTC-A
216HCV-TR3bTGTGC-TC-A
217NE483cATAC
218NE2743dGTGCCCTGGCCTCC-
219NE1453eATGC
220NE1253fATAC-TCCTAGCCCT-A
221Z44aTATGTCACTAT-A
222Z14bTTA-CCAA
223GB3584cTACGCAA-C-A
224DK134dTCGCACA
225GB8094eTACCGTGCAA-C-A
29BNL74kT-TC-A
31BNL84kCA-T-TC-A
33BNL94kTTACCGTACA
35BNL104kT
37BNL114k
39BNL1241
226BE955aTTTATTCCACTA-A
227HK26aTCACCCCCTGA
228FR17aTC-TAACCCT-TTA
43VN48aTCA
47VN129aTTC-ACCTAGCCAA
51NE9810aATGATCCAGGGTCTC-G



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Fig.1N

SEQID	
NO651700	
	ACACTCCGGGGTGCGTCCCTTGCGTTCGTGAGGGCA
	-TCTGCTGC-AAC
	ACAGC
	-CATGCGAA
11FR21fGCATT	-GTNGCA-AGA
	GTCCGAGAAA-TG-
	-TCTT-AA-T-AGAATAATG
	T-AT-AGACC-C
	-GTCCT-AGGAGA
	GTTATCAGAA-AGCTC-G TTATCT-AGA-GTCAT-
	GTTATGT-AGA-G1CA1-
	GTCTATTAT-AGA-GC-CCAA-
	GTCTATTCT-AGT-GAA
215NZL13aTTT	ACCTATC-AGC
216HCV-TR3bATGT	TTACAGCCACAACC
	-TTGCTACC-AAA-CAAT-
	TTGATTGCAATCA
	TGTTTCG-AGA-C
220NE1253fTAT	TGCTGCACCT-
221744- 2 22 2	ma , a m aymayam a
	TGG- TGATCTGGACAG-
	CTCATT-ACGA-G-TTG-
	-CTCATGA-GAG-
	CTCAACTGAAGACCG-
	CTCATGCGA-AG-
31BNL84k-C-CCAT	CTATGCGA-AACTG-
33BNL94k-CCAT	CTCATGCGA-A-TG-
	ACTATGCGA-A-TG-
	-CTAAA-
39BNL1241-CCAT-A	-CTAATACTGAAGACTG-
22600055-#3 00#0 3	C 3
226BE955aTA-CCTGA	-G-ATTGTCATGACAT-
227HK26aT-C-ATGTT	TTGTAT-GTGA-G-TC-ATG
ZZ/MKZOGI C AIG I I	TIO I A I O I GA G IC-AIG
228FR17aGACCATGAT	CTATTATA-CAAG-CG-
43VN48aGACACTGTTT	TGTTATGAAGRT-RA
47VN129aT-GCATGT	CTCTCGAAGACC
EINEGGIOGG AMM C	
SINESSIUAGATTC	-TTATCTCACTCT



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Fig.10

```
SEQID
NO701750
208HCV-11aACGCCTCGAGGTGTTGGGTGGCGATGACCCCTACGGTGGCCACCAGGGAT
209HCV-J1b-TTT---CC-T--C---A---C-C--T--C--C--GG----A-C
3BNL11d--CATCTCC-C--C--A----C-C----C-T--GGT--AAA-Y
7BNL21d--T-T--TC-T--C---A--C-RC-C----C-T--GGT--AA--C
11FR21f-TAT---CC-T--C----AC--C-C--C--C--C--AG-GC--ATC
211HC-J62a-TA-A--TC----C---A-AC--G-CT-A--G-AT------GTGCA-C-G
212HC-J82bG-A---T-CAT--C---A-ACAAG-A--A--C-AC-----TGTG-AAC-C
213S832c---T---TC-A------C--G-TG----C-ATC-C----TA--TC-A
214NE922d--ATA--CC-C-----A-AC--G-TT-G--C-ATA-A--TGTG--CC-A
15BNL32eGTCGG-TCCAC----A-CC---CT-G--C-ACA-A---GTG--CA-A
17FR42f-TAGGA-CTTC-----ACA---G-CT-G--C-AC----TGTG--CCGA
19BNL42g-TAAG--CC----C--A-AC--G-C--T--C-AC----TGTG-ACC-G
23BNL52h-TCAG--TC-C--C--A-AC-TG----A--C-AT-----GTG--CC-A
25BNL62i--A----CC-C--C--A-AC--G-C-----ACA-C--TGTG--CC-A
215NZL13a-TA-A--T-C---C---ACCC-AG----A----A----AGT----T-C
216HCV-TR3b--CAA--ATCA--C---ACAA--G-CT-AA-G------GTT---ACC
217NE483c--A--A---C---C---A-AC--G----T--G--A-----GGT---TC-C
218NE2743d--T----CAA--C---A-TC--G--G-A--A--A--A-----GGTT-A-T-C
219NE1453e--A-A---GA---C---ACCC--GC---A--A------AGT---AT-C
220NE1253f--CAG--A----C--AC-C-AG-A--A--G--A----TGT--AAC--
221Z44a--A-A---C-T--C---AC-C--G----G----A----TGT-GCAC-C
223GB3584c-TCAG--AC-C--C------CC-C--T--C--C-----GG-GCCTT-C
224DK134d--AAG--T-CA--C----T-TC-C-----C--C---TG-GCAAC--
29BNL74k-TCAG--AC-T--C----A--CC-T------C--C--AG-GCCAT-C
31BNL84k-TCAG--AC-T--C------CC-T--T-----C--C--AG-GCCAT-C
33BNL94k-TCAG----T--C------CC-T------CA-C--AG-GCCAT-C
35BNL104k--CAG--AC-C--C------CC-T------C--C--AG-GCCAT-C
37BNL114k-TCAT--AC-C--C------CC-T-----C--C--AG-GCCAT-C
39BNL1241--A-T---C-C--C------CT-A--A------G-GCCCATA
226BE955a-T-TGAGT--A--C----CCAA--T-----AC--T-AG--CC-AGC
227HK26a-TCGG--C-CC-----CAT--TG-----C-CC-----TACCAA--
228FR17a-T-AG--AC-A----C-CC-TG-CT----C-CT-A---GT-CCCA-C
43VN48a-TCAA--CC----C----CA-GCCT----G--CC----AGTGCC-A-C
47VN129a--CTGA-C-A----C--T--GCCT----G--AT----GGTGCA-A--
51NE9810a-TA-A--A--A--C--A-CC-TG---G---Y--C--C---GTG-A-TCG
```



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Fig.1P

```
SEQID
NO751800
208HCV-11aGGCAAACTCCCCGCGACGCAGCTTCGACGTCACATCGATCTGCTTGTCGG
209HCV-J1bA---GCA----A-C--ACAA-A----C--G----T---C--T--
210HC-G91cTCGCGCG-----TC-GTG--G----GTG----CTC-A-----
3BNL11d-CT-GTG----A-TR--GCAA-C-----G---CT-----T--
7BNL21d-CT--TG----TA-TG--GCAA-C----C--TG----CT----G--T--
11FR21f-CG--CGCT---ATCGATG--G-G--G-----G----C--C--C--G--
211HC-J62aCC-GGCGC--T-A--CA-GGCT-A--GACG-----T--CA--G----GAT
212HC-J82bC--GGTGCG-T-A-TCGTAGC--G---ACA---G---CA--A-C--AAT
213S832cCCTGGCGCT-T-A-T-A-GGC--G---GCA------A-CA-C--GAT
214NE922dCCTGGTGCG-TTA-C-A-GGC--G--GACG--T--T--ACCA-CA-T-C
15BNL32eCCTGGTGCT-T-A-C-A-GGA--G--GGCA-G---T---GCCG-C--GAT
17FR42fCCTGGTGCT-T-A-T-GAGGT--G--GGC-----T---ACCA-C--GAT
19BNL42qCC-GGCGC-T-A-T-G-GGCT-G--GACG----T--CACCA-C--GAT
25BNL62iCCTGGCGCG-TTA-C-A-GGC--G--GACA--T--T--CA-CA-----C
215NZL13a-T-GG-GCAA-TA-TG-TTC-A-A--CA----TG-G--C--AT-A--A--
216HCV-TR3bCTTGGCG-GA--A-CG--TC-A-C---ACC--TG-G---A---G--A--
217NE483c-T-GGTGCGA--A-CG-ATC-A-C--CG-G---G-G--------G--G--
218NE2743d-CTGGCGCGA--A-TG-ATC-A-C--CA----TG-G-------G--G--
219NE1453e-CTGGTGCAA-GA-G-TTCCG-A--CG-A--G-G--T----A----
220NE1253fCCTGGCGCAGT-A-CG-ATCAA-C--CA-G--TG-G---T--A-G--G--
221Z44aCCGGGCGCT--GCTTGA-TC-T-C--G--A--TG-G--CT-AA-G--A--
222Z14bCC---CGCA--GTTAGA-TCCA-G--CA-G--TG-A--C--A-G--G--
223GB3584cAT-GGCGCT--GCTTGAATCC--C--GA----TG-G-----A-G--A--
224DK134dCTG--TGCT--GCTTGA-TCTT-GA------G-G-----A-G--G-
225GB8094e-T-GGTGCT--GCTCGA--CCT-G--G--C--TG-G--C--A-G--A--
29BNL74kAT-GGCGCG--ACTTGA-TCT--A--GA----TG-G--CT--A-G--G--
31BNL84kAT-GGCGCA--GCTTGA-TCT--G--GA----TG-G-----A-G--G--
33BNL94kAT-GGCGCA--GCTTGA-TCCT-G--GA----TG-G-----A-G--G--
35BNL104kAC-GCGGCG--GCTTGA-TCC--G--GA----TG-G-----A-G--G--
37BNL114kAT-GGCGCG--ACTTGA-TCT--A--GA----TG-G---G--A-G--G-
39BNL1241CTTTCGGCT--ACTT-T-TCCG-A--G--G--TG-G-----A-G--G--
226BE955aCT-GG-GCAGT-A--G-T-CT----GA-AGC-G-T--CTAC--A-CG-
227HK26a-CTTCCACG----A---GGAT-C--CA-G--TG-G----T---CG--
228FR17aTCATC-G-G--AATCCACGG-T----C-A---G-A--C--C--T--
43VN48a-CGTCTACG--A-TC--CGG-T-C--CAAA--TG-G--CA-CA-G--G--
47VN129a-CGTCGG-GT--ATC-G-GGTG-C--CGAG---G-G--C--CT-G--G-
51NE9810aCC-TGCGC-G--A-CG-CTCT--C--CACG---G-G--A--A-G--G--
```



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Fig.1Q

SEQID NO801850 208HCV-11aGAGCGCCACCCTCTGTTCGGCCCTCTACGTGGGGGACCTATGCGGGTCTG 209HCV-J1b-GCGTG-T
211HC-J62a-TCGCCTTCTGGG- 212HC-J82b-GCATGGCCT-GTATG-GG-C- 213S832c-TCTTGGTTT
15BNL32e-TCCTGATG-GCG-A- 17FR42f-TCCG-T
19BNL42g-GTTGTAA-CG-GTCG-G- 23BNL52h-TCTTGCATT-GTCT-CCG-A- 25BNL62i-TCGTCTT-GT
215NZL13aCGCGGA-GCTGTTA-GTG 216HCV-TR3bCGCACGACAAGGGCGCT-TG
217NE483cT-CGTAT-GATCTG-A- 218NE2743dAGCTTGT-GCCGGTTCTA-GTAG-C-
219NE1453eCTTGCCGTCTTG-C- 220NE1253fTGCAGGAATTATT-GG
221Z44aCGCGTT-GTTTTCAGG 222Z14bTGCGTTA-GCTA-TA-TGTAGGC-
223GB3584cTGCTTGCGCCTTATGTGGC- 224DK134dCGTCCCA-CAGTGG-
225GB8094eTGCTG-GCCCCCTGGCT 29BNL74k-GCTG-TATA-CTT-RTYGGCT
31BNL84k-GCTTG-TC-ATA-CTT-GTCGGCT 33BNL94k-GCGTGATA-CTT-GTCGG 35BNL104kAGCTTG-TATA-CYT-GTCGGCT
37BNL114k-GCTTG-TATA-CT-GGCT 37BNL114k-GCTTG-TATA-CT-GGCT 39BNL1241TGCATA-CGTT-ACGG
226BE955aAG-GTGCCGT-AAAGCGTG-AC 227HK26aCGCAGTGG-TCATG-A-C
228FR17a-GCAGG-AT-TA-GA-CA-CTTAGCA
43VN48aCGCTG-GTATA-GTGCCC
47VN129aTGCTTG-GTCTA-GCTTGGGC
51NE9810aRGCGACATAATTAG-GC



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Fig.1R

SEQID
NO851900
208HCV-11aTCTTTCTTGTCGGCCAACTGTTCACCTTCTCTCCCAGGCGCCACTGGACG
209HCV-J1b-TCTCGATC-CGT-TGA
210HC-G91cCTGA-CAT
3BNL11dCC-CTGATAC-CATGCATA 7BNL21dC
11FR21fCTGTA-GTCGT
11FR211CC1G1
211HC-J62a-GA-GCA-CGATTGGACAATTT
212HC-J82b-GA-GAC-ATCGGGCTTGG-AA-ACAAAACTTC
213S832c-GA-GG-CCTGG-CGGT-G-GGACAA-ATAC-TTT
214NE922d-GA-GT-G-CTTCTG-CT-AGCAATTAA-TTT
15BNL32e-GA-GA-A-CT-CAGGCTT-G-GG-AG-AT-ACTTC
17FR42f-GA-GA-A-CA-CGG-TGC-GT-GAGCAATATACTTTT
19BNL42g-GA-GA-A-CT-CTGG-TGTTGGGCAA-ATAACTTT
23BNL52h-GA-GT-GTCTT-TTGACTCAAATCTTC
215NZL13aCGAGCCGAGATC-ATCAA
216HCV-TR3b-GGAGCAGATC-CACC
217NE483c-TCCAAGCAAAGAC-ACAA
218NE2743dCT-GGAGGCTAGATC-T-AGAAC
219NE1453eCGGGGCCTAAGGTC-TTACT
220NE1253f-TCGTAGAG-TCAAT-ATC
221Z44aCCGA-GGGAATTCGGGC-TC
222Z14bCAGGGACGAGC-CGC
223GB3584c-AT-GTTGAT-TCAGGCT
224DK134d-GCT-GT
225GB8094e-ACT-GAACAAGC-A
29BNL74k-GCATGAT-TCGAAT
31BNL84k-GCT-GTTGATT-TCGAAC-AT
33BNL94kCGCT-GTTGAT-TCGAAC
35BNL104k-GCT-GTTGAT
37BNL114k-GCGTTGAT-TCGAAC
39BNL1241CCAGGA
226BE955a-ACT-GAAATAGGTC-C-AGGCT
227HK26aT-G-CGAATCAGC-CTT
228FR17a-AA-CT-GAGGTTTAGGT-A-TATCA-GTT
43VN48a-TCCTAGCGCAGGTCATGTCA-GTT
47VN129aCTGGTGAGAATGT-TGATC
51NE9810a-AYGGGGT-A-GGAGA-ATC-C-AGTT



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Fig.1S

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SEOID
208HCV-11aACGCAAGGTTGCAATTGCTCTATCTATCCCGGCCATATAACGGGTCACCG
209HCV-J1bGTA----A------A-----A------CG--T-A-----
3BNL11d----G-AG----C---A---
7BNL21d--A--G-AG-----C----A---
11FR21fGT---G-AC--T----T--C--T--CT-T-----C------
211HC-J62aGT----AC-----C--T-TACC--C--T--A----
212HC-J82b--C----AG-----C--T--C--AA--T--C--C--C--T--
213S832cGTC--G-AA-----C--T--C--A--C--GC--T-----A----
214NE922dGTC--G-AC-----C--T--C--A--C--A-----C--C--T--A--T--
15BNL32eGTC--G-AA------C--A--C--A-----C--T--A----T--
17FR42fGTC--G-AA-----C--A--C--A--C--A------C--A--T--
19BNL42gT-C--G-A-----T--C---
23BNL52hGTC--G-A----C----G--A
215NZL13aGTC--GACC--T--C----GC-G--C--A-----C-TT-A--A--T--
216HCV-TR3bGT---GACG-----C----G--A--C--A------G-TT-A--A--T--
217NE483cGTT--GCA-----C----AC-G--C--A--T---G-TT-A-----T--
218NE2743dGT---GACC------AC-G--C--T--T--C---T-A--A--A-
220NE1253fGTC--GTTG------AC-A--C--A--A--C--T--A--A--T-A
221Z44a----G-AG------T--C-----CA-T-------C--C--C--A-
222Z14b--C--G-A-----C----C----T--T--CG-CT----C--A-
223GB3584c----G-AC-----T--C-----CG-G--G--CG-T-----C--A-
224DK134d--C---AC-----T--C-----CA-A--A-----C--A--A--A-
225GB8094e--C--G-AC--T-----T--C-----CG-A--G-----T------C--T--
29BNL74k--T---A-----T--C---
31BNL84kG-C--G-A-----T-----
33BNL94k--C---A----C---
35BNL104k--C--G-A-----T--C---
37BNL114k--C--G-AA----T--C---
39BNL1241GTC----AC-----C--T--C---
226BE955aGT---GAAC----C-T--C-T--CAGT-----G-T--C--C----
227HK26aGT----AC----C----A-A----CG-C--C--A-
228FR17a--C--G-A---T--C------NA-CN-T-----CG-C-----A---A-
43VN48aGTC--G-AG--T--C--T--C-----CA-A--G-----C--T--A----
47VN129aG-C--G-AC----C--T--C-----G-A----C--C--T--G----
51NE9810aGTC--G-AC----C--T--C---
```



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Fig.1T

SEQID NO951957 208HCV-11aCATGGCA 209HCV-J1bT 210HC-G91cAT 11FR21fNNNNNNN
211HC-J62aG 212HC-J82b 213S832cT 214NE922dGG 15BNL32eG 17FR42fANN
215NZL13aAT 216HCV-TR3bTG 217NE483cGT 218NE2743dGT 219NE1453e
221Z44aGG 222Z14bGC 223GB3584cG 224DK134dAT 225GB8094eGT
226BE955aG 227HK26aGT
228FR17aG
43VN48aA 47VN129aGG



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Fig.2A

SEQID NO150
229HCV11aMSTNPKPQKKNKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATR
230HCV-J1bR-T
2BNL11dX-TXXXXXXX
6BNL21dR-TXX
10CAM10781eA
12FR21fR-T
231HCJ62aR-T
232HCJ82bR-T
233CH6102cR-T
234NE922dR-T
14BNL32eR-TR-T
18FR42fP-
235HCVTR3bLRQTLNV-
236DK134dR-TMM
237CAM6004eR-TMM
238GB8094eL-R-TMM
28BNL74kR-TM
239BE955aR-TM
240HK26aLR-TT
42FR17aLR-TM
44VN48aLR-TI
46VN138bLR-T
48VN129aLR-TM
50NE9810aLR-TXVQV-



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Fig.2B

SEQID N051100 229HCV11aKTSERSQPRGRRQPIPKARRPEGRTWAQPGYPWPLYGNEGCGWAGWLLSP 230HCV-J1b
231HCJ62a
235HCVTR3bKQ-HLSRSKL
236DK134d
238GB8094e
28BNL74kX
239BE955aALQ-TS-GAL
240HK26aQ-QH
42FR17a
44VN48a
48VN129aA
EONEGO 100 C D T C



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Fig.2C

SEQID
NO101150
229HCV11aRGSRPSWGPTDPRRRSRNLGKVIDTLTCGFADLMGYIPLVGAPLGGAARA
230HCV-J1b
2BNL11dN
6BNL21d
12FR21fS-T
231HC-J62aVVVVV
232HC-J82bTHRIVVV
233CH6102c
14BNL32eXXX-VVX
18FR42fVVVVVVV
101.421
235HCV-TR3bVVV
241GB1164cVV
236DK134dVVV
237CAM6004e-XXNX
238GB8094eVV
242G224fVV
243GB5494gVV
244GB4384hVV
28BNL74kN
239BE955aNK
237DD333dKK
240HK26aVV-A-
42FR17aNNXXLVL-GV-A-
44VN48aVXV-X-
46VN138bXNXXXIE
40101100- D V N V D D W N N N N N N N N N N N N N N N N N
48VN129aD-X-NX
50NE9810aN



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Fig.2D

	SEQID
	NO151200
	229HCV11aLAHGVRVLEDGVNYATGNLPGCSFSIFLLALLSCLTVPASAYQVRNSTGL 230HCV-J1b
	2BNL11dXT-HEAS-V
	6BNL21dFTT-HEAS-V
	12FR21f-XXGXXXXXXXXXTE-HST-DG
	ILIKETI K NO MMAKA K KA K I E NOI DO
	231HC-J62aFFI-T-VAE-K-ISTG
	232HC-J82bIISSS
	233CH6102cISISVVE-K-TSTS
	234NE922dIIIV-GLK-TSSS
	14BNL32eXIXXVV-XVE-K-TSQA
	18FR42fIK-NSHF
-	- କଥାଯାଁ L42g
	24BNL52hIVK-TSHS
	26BNL62iIIVVA-RS-S
	00511011 8701
	235HCV-TR3bA-GFFCGLEYT-TS
	241GB1164c-EAVI
	236DK134dLNYS-V
	237CAM6004eAVITVNYAS-I
	238GB8094eAVI
	242G224fAVI
	243GB5494qAVI
	244GB4384hAVI
	30BNL74kI-FINYVS-I
	32BNL84kIINYTS-I
	34BNL94kIINYH-TS-I
	245BNL94kIIXXTNYVS-I
	36BNL104kIXTNYVS-I
	38BNL1141IIQHYVS-I
	239BE955aVPYAS-I
	237DB7554VF1A5-1
	240HK26aAIITLTYGS
	42FR17aAI
	44VN48aXXIXXXX-XXXTAHYT-KS
	TATALAN MATANANAN ANAN ANAN ANAN ANAN ANAN AN
	48VN129a-XAIIXTLNYA-KS
	52NE9810aI-F



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Fig.2E

		_
SEQID		
CTATA		
370001000		
NO201250		

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NO<sub>2</sub>
229HCV-11aYHVTNDCPNSSIVYEAADAILHTPGCVPCVREGNASRCWVAMTPTVATRD
230HCV-J1b-----S-------M-M-----S-F-----L--L-A-N
2BNL11d-----S----I--MDGM-M-Y-----D-HL---M-L---L-VKX
6BNL21d--L---S----I--MSGM---A-----N-S----MXL---L-VK-
12FR21f-----S-G-----K-I----X---I----PL---L-A-I
231HC-J62a-M----T-D--TWQLQA-V--V-----EKV--T----IPVS-N--VQQ
232HC-J82b-YA----S-N--TWQLT--V--L-----ENDNGTLH--IQV--N--VKH
233CH6102c-M----S-----WQLEG-V------EQI------PVS-N--I-Q
234NE922d-M----Q-----WQLR--V--V-----EEK--I----IPVS-NI-VSQ
14BNL32e-MA----S-N---WQLX--V--V-----ENSSGRFH--IPIS-NI-VSK
18FR42f-MA----A-D---WQLR--V--V----E-S--RTF--T-VS-N--VSR
20BNL42g-MA----S-N--IWQMQG-V--V-----ELQ--K----IPV--N--VNQ
24BNL52h-M----S----WQLK--V--V----E-HQ-Q----IPV--N--VSQ
26BNL62i-M----S----WQLEE-V--V-----EWKD-T----IPV--NI-VSQ
235HCVTR3b-VL---S-G-----E-V---L----TT--O-S--TTVST---V-T
236DK134d------TDYH---L-----K-T---SL----AQH
237CAM6004e--I----A----TENH---L-----T--Q-----L----SPY
238GB8094e--I-----A----TDNH---L-----KT--Q-----L----SPY
242G224f--L-----F--VHH---L-----T--Q-----L--L-APY
30BNL74k-Y-----L-----DHH---L-----Q-----L-----APY
32BNL84k------DHH---L-----T-Q-----L----APY
245BNL94k-----L-----DHH--AL------V--Q------L-----APY
36BNL104k-----L----F--DHH---L------K---H----L-----APY
38BNL1141-----L----SDHH---L-----KT--T-----L----API
246GB7244x--I-----V---TDHH---L----T--V----TPV-----AVS
239BE955a-----QI---LSAPS
42FR17a--L---S-N---F--ETM---L-----IKA--E----LPVS--L-VPN
44VN48a--L------ETL---L----KXX-Q-----QAS--L-VPN
48VN129a--L------NGM---L-----KT--LTK--LSAS--L-VQN
52NE9810a-M----S-G-----G-I---L-----S-T----IPVSX---VKS
```



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Fig.2F

SEQID NO251300 229HCV-11aGKLPATQLRRHIDLLVGSATLCSALYVGDLCGSVFLVGQLFTFSPRRHWT 230HCV-J1bSSI-T-TIVA-AMSYE- 2BNL11dASV-TXAIVXX-FM-XA
231HC-J62aPGALTQG-TMV-M
235HCVTR3bLGVTTASI-T-V-MARQAF-AAF-AT-
241GB1164cVGA-LESS-V-M-A-VIGM-S-Q 236DK134dLNA-LESV-M-GI-V-GQ 237CAM6004eAGA-LEPV-M-A-MIGLMQ 238GB8094eVGA-LEPV-M-A-V
239BE955aLGAVTAPAV-Y-A-G-AAALMYRQ-A-
240HK26aASTGFVA-A-VVSILAQ
42FR17aSSV-IHGFVA-AFM-IIIR-KY-QV
44VN48aAST-V-GF-K-V-IMA-AFMGLLRM-QV
48VN129aASVSIRGV-E-VA-AFMGLRMYEI
52NE9810aPCAATAST-V-MM-XAALXG-SWRH-Q



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Fig.2G

SEQID
NO301319
229HCV-11aTQGCNCSIYPGHITGHRMA
230HCV-J1bV-DVS
2BNL11dE
6BNL21dE
12FR21fV-DSXXX
231HC-J62aV-DT
232HC-J82bEQ
233CH6102cV-EX
234NE922dV-D
14BNL32eV-E
18FR42fV-EX
20BNL42gS-D
24BNL52hV-D
235HCVTR3bV-TVS
241GB1164cDAV
236DK134dDT 237CAM6004eDT
237CAM6004eDT
238GB8094eDA
242G224fET
243GB5494gDD
244GB4384hDV
30BNL74kD
32BNL84kA-D
34BNL94kD
245BNL94kD
36BNL104kE
38BNL1141V-D
246GB7244xDT
239BE955aV-NSV
240HK26aV-DTV
42FR17aDXNXV
44VN48aV-ET
48VN129aA-DA
52NE9810aV-D



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Fig.3A

SEQ ID NO. 1 (BNL1, 1d)
ATGAGCACGAATCCTAAACCTCAAAGAAAAACCAAACGTAACACCAACCGCCGCCCTCAKGGSGTN
NNNNNNCCGGGTGGCGGTCAGATCGTTGGTGGAGGTTTACCTGTTGCCGCGCAGGGGCCCCAGGNNG
GGTGTGCGCGCGACTAGGAAGACTTCCGAGCGGTCACAACCTCGTGGCAGCCGACAGCCTATCCCC
AAGGCTCGYCGGYCGAGGGCAGGTCCTGGGCTCAGCCCGGGTATCCTTGGCCCTCTATGGCAAT
GAGGGCTGCGGGTGGCCGGGNTGGCTCCTGTCCCCCCGCGGCTCTCTGGCCCCAATTGGGGCCCC

SEQ ID NO. 3 (BNL1, 1d)
GACGGCGTGAACTATGCAACAGGGAACTTGCCCGGTTGCTCTTTCTCTATCTTCCTCTTGGCTTTG
CTGTCCTGCTTGACGGTTCCAACKACCGCTCACGAGGTGCGCAACGCATCCGGGGTGTATCATGTC
ACCAACGACTGTTCCAACTCGAGGATCATCTATGAGATGGACGGTATGATCATGACTACCACGGG
TSCGTGCCCTGCGTTCGGGAGGATAACCATCTCCGGCTGGATGGCGCTCACCCCCCACGCTTGCG
GTCAAAAAYGCTAGTGTCCCCACTRCGGCAATCCGACGTCACCTTGCTTGTTGGGGGNNCC
ACGTTCTGTTCCGCCTATGTACGTGGGRGACCTTTGCGGGTCTGTCTTCCTCGCTGGCCAGCTATTC
ACCTTTTCACCCCGCATGCACCATACAACGCAGGAGTGCAACTGCTCAATC

SEQ ID NO. 5 (BNL2, 1d)
ATGAGCACGAATCCTAAACCTCAAAGAAAAACCAAACGTAACACCAACCGCCGCCCACAGGACGTC
AAGNTCCCGGGTGGTGGTCAGATCGTTGGTGGAGGTTTACCTGTTGCCGCGCAGGGGCCCCAGGTTG
GGTGTGCGCGGCACCAGGAAGACTTCCGAGCGGTCGCAGCCTCGTGACAGGCGACAGCCTATTCCT
AAGGCTCGCCAGTCCGATGGCAGNNCCTGGGCTCAGCCAGGCATCCCTGGCCCCTCTATGGCAAT
GAGGCTGCGGATGGCGGGGATGGCTCCTGTCCCCCCGCGGCTCTCGGCCCCAGTTGGGGCCCC

SEQ ID NO. 7 (BNL2, 1d)
GACGGCGTGAACTATGCAACAGGGAATTTGCCTGGTTGCTCTTTCTCTCTTTAGCTTTT
CTGTCCTGCTTGACGGTTCCAACTACCGCTCATGAGGTGCGCAACGCATCCGGGGTATATCATCTC
ACCAATGACTGTTCCAACTCGAGCATCATCTATGAGATGAGTGGTATGATCTTGCACGCCCCAGGG
TGTGTGCCCTGCGTTCGGGGAGAACACTCTTCTCGTTGCTGGTTGCACTCTCACCCCCACGCTTGCG
GTCAAAGACGCTAATGTCCCTACTGCGGCAATCCGACGCCATGTCGACTTGCTGGTTGGGACAGCC
GCGTTTCGGTTCCGCTATGTACCTGGGGGACCTCTGCGGATCCGTCTTCCTTGTCGGCCAGCTATTC
ACCTTTTCACCCCGCTTGTACCATACAACACAGGAGTGCAACTGCTCAATC

SEQ ID NO. 9 (CAM1078, 1e)
ATGAGCACGAATCCTAAACCTCAAAGAAAAACCAAAAGAAACACCAACCGCCGCCCACAGGACGTC
AAGTTCCCGGGCGGTGGCCAGATCGTTGGTGGAGGTCTACGTGCTACCGCGCAGGGGCCCTAGATTG
GGTGTGCGCGCAGCGCGGAAGACTTCGGAGCGGTCGCAACCTCGTGGGAGGCGCCAACCTATTCCC
AAGGAGCGCCCACCCGAGGGCAGGT



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Fig.3B

SEO ID NO. 11 (FR2, 1f) ATGAGCACGAATCCTAAACCTCAAAGAAAAACCAAACGCAACACCAACCGCCGCCCACAGGACGTT AAATTCCCGGGTGGGGGCAGATCGTGGGTGGAGTTTACTTGTTGCCGCGCAGGGGCCCCAGGTTG GGTGTGCGCGCGACGAGGAAGACTTCCGAGCGGTCGCAACCTCGCGGAAGGC GACAGCCTATCCCCAAGGCTCGCCGACCCGAGGGCAGGTCCTGGGCTCAGCCTGGGTACC CATGGCCCCTCTATGCTAACGAGGGCTGCGGATGGGCGGGATGGCTCCTGTCCCCTCGCG GCTCCCGTCCTAGCTGGGGCCCCAATGACCCCCGACGTAGATCACGCAATTTGGGTAAGG TCATCGATACCCTAACGTGTGGCTTCGCCGATCTCATGGGGTACATTCCGCTCGTCGGCGC CCCCTAGGGGGCGCTTCCAGAACCCTGNCACATGGTGTCCGGGTCCTGGNAGGCGGCGTGATNNN NNNNNNNNAACCTTCCNGGTTGCTCTTTNNCTATCTTCCTCTTGGCNTTACTCTCTTGCCTCAC AGTCCCCACCTCTGCCTATGAGGTGCACAGCACAACCGATGGCTACCATGTCACTAATGACTGTTC CAACGGCAGCATCGTATATGAGGCAAAGGACATCATCCTTCACACGCCTGGGTGNGTGCCCTGCAT ACGGGAAGGCAATATCTCCCGTTGCTGGGTACCGCTCACCCCCACGCTCGCAGCGCGGATCGCGAA CGCTCCCATCGATGAGGTGCGGCGTCACGTCGACCTCCTCGTGGGGGCAGCCGTGTTCTGCTCAGC CATGTACATTGGGGACCTTTGTGGGGGGCGTCTTCCTCGTTGGGCAATTGTTCACCTTCACGTCCCG GCGGCATTGGACGGTGCAGGACTGTAATTGTTCCATTTACTCTGGCCACATAACGGGCCACCGNNN NNNN

SEQ ID NO. 13 (BNL3, 2e)
ATGAGCACAAATCCTAAACCTCAAAGAAAAACCAAAAGAAATACCAACCGCCGCCCACAGGACGTC
AAGTTCCCGGGCGGCGGCCAGATCGTTGGCGGAGTTTACTTGTTGCCGCGCAGGGGCCCCAGATTG
GGTGTGCGCGCGACGAGAAAGACTTCTGAACGGTCCCAGCCACGTGGAAGGCGCCAGCCCATCCCT
AAAGATCGGNGNGCCACTGGCAGGTCCTGGGGACGTCCAGGATATCCCTGGCCCCTGTATGGGAAC
GAGGGGCTCGGCTGGGCAGGATGGCTCCTGTCCCCCCGAGGCTCTC

SEQ ID NO. 17 (FR4, 2f) ATGAGCACAAATCCTAAACCTCAAAGAAAAACTAAAAGAAACACTAACCGTCGCCCACAGGAC GTTAAGTTCCCGGGCGGCGGCCAGATCGTTGGCGGAGTTTACTTGTTGCCGCGCAGGGGCCCCAG GTTGGGTGTGCGCGCCCAAGGAAGACTTCTGAACGGTCCCAGCCACGTGGAAGGCGCCAGCCC ATCCCAAAAGATCGGCGCGCCACTGGCAAGTCCTGGGGACGTCCAGGATACCCTTGGCCCCTGT ACGGGAACGAGGGCTCGGCTGGCAGGGTGGCTCCTGTCCCCCGGGGGCTCTCGCCCCTCGTG GGGCCCAAACGACCCCCGGCACAGGTCACGCAACTTGGGTAAGGTCATCGATACCCTCACGTG TGGCTTTGSCGACCTCATGGGGTACATACCTGTCGTCGGCGCCCCTGTGGGCGGCGTTGCCAGA GCCCTCGCGCATGGCGTGCGGGTCCTGGAGGACGGGATAAATTATGCAACAGGGAACTTGCCCGGT GTTAAGAACAACAGCCACTTCTACATGGCGACTAATGACTGTGCCAATGACAGCATCGTCTGGCAG CTCAGGGACGCGGTGCTCCATGTTCCTGGATGTGTCCCCTGTGAGAGGTCAGGTAATAGGACCTTC TGTTGGACAGCGGTCTCGCCCAACGTGGCTGTGAGCCGACCTGGTGCTCTCACTAGAGGTCTGCGG GCTCACATTGATACCATCGTGATGTCCGCCACCCTCTGCTCTGCCCTATACATAGGGGACCTATGC GGCGCTGTGATGATAGCAGCGCAAGTTGCCGTCGTCTCACCGCAATACCATACTTTTGTCCAGGAA TGCAACTGCTCCATATACCCAGGCCATATCACAGGACATCGAATGGNN



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Fig.3C

SEQ ID NO. 19 (BNL4, 2g)
GACGGGGTAAATTATGCAACAGGGAATCTGCCTGGTTGCTCTTTCTCTATCTTCTTGTTGGCTCTT
GACGGGGTAAATTATGCAACAGGGAATCTGCCTGGTGAAGAACACCAGTACCATGTACATGGCA
ACCAATGACTGTTCCAACAACAGCATCATCTGGCAAATGCAGGGCGGGTGCTTCATGTTCCTGGA
ACCAATGACTGTTCCAACAACAGCATCATCTGGCAAATGCAGGGCAATACCGGTCACTCCCAACGTGGCT
TGTGTCCCGTGTGAGTTGCAGGGCAATAAGTCCCGGTACACCATCGTGATGGTCGCT
GTGAACCAGCCCGGCGCCCTCACTAGGGGCTTGCGGACGCACATTGACACCATCGTGATGGTCGCT
ACGCTCTGTTCTGCACTCTACATCGGGGACGTGTGTGCGATTGTTCCATC
ATTGTCTCGCCGCAACATCACAACTTTTCCCAGGATTGCAATTGTTCCATC

SEQ ID NO. 21 (BNL5, 2h)
ATGAGCACAPATCCTARACCTCARAGARARACCARAGARACACTARCCGCCGCCCACAGGACGTT
ATGAGCACARATCCTARACCTCARAGARARACCARAGARACACTTACTGCCGCGCAGGGGCCCCCGGTTG
ARGTTCCCGGGCGGTGGCCAGATCCTTGGCGACGCCACCCAGCCCATCCCT
GGTGTGCGCGCGACGAGGARACCTTCCGAACGGTCCAGCACCTTGGCCCCTGTATGGGARC
ARAGATCGGCGCTCCACTGGCARATCCTGGGGACGTCCAGGATACCCTTGGCCCCTGTATGGGARC
GAGGGCCTTGGTTGGGCAGGATGCTCTTGTCCCCTCGAGGCTCTC

SEQ ID NO. 27 (BNL7, 4k)
ATGAGCACGAATCCTAAACCAAAGAAAAACCAAACGTAACACCAACCGCCCCCATGGACGTT
AAGTTCCCGGGTGGTGGCCAGATCGTTGGCGGAGGTTTACTTGTTGCCGCGCAGGGGCCCCAGGTTG
AAGTTCCCGGGACTCGGAAGACTTCGGAGCGGTCGCAACCTATCCCC
GGTGTGCGCGCGACCTCGGAAGACTTCGGAGCGCCCAGCCTATCCCCCAACCTCTCGGAGCGCCGAACCTTCCGCAACCTCTCGGGAAGACTCCTGGGCACCAGCCCAGGATATCCATGGCCTCTTTACGGTAAT
AAGGCGCGTCGATCCGAGGGAAGGTCCTTGTCCCCCCGCGGTTCTC
GAGGGTTGCGGGCGANNATGGCTCTTGTCCCCCCCGCGGTTCTC



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Fig.3D

SEQ ID NO. 35 (BNL10, 4k)
GACGGGATCAATTATGCAACAGGGAATATTCCCGGTTGCTCYTTTTCTATCTTCCTTYTGGCACTT
CTCTCGTGTCTGACTGTCCCCGCTTCGGCCACTAACTATCGCAACGTCTCGGGCATCTACCATGTC
ACCAATGACTGCCCGAATTCAAGCATAGTGTATGAGGCCGACCATCACATCTTAGCACTTCCAGGT
TGCGTGCCCTGCGTGAGAGTGGGGAACCAGTCACGCTGCTGGGTGGCCCTTACCCCTACCGTCGCA
GCGCCATACACCGCGGCGCCGCTTGAGTCCCTGCGGAGTCATGTGGATCTGATGGTGGGAGCTGCC
ACTGTTTGTTCAGCCCTTTACATCGGGGAYTTGTTGTGGCGGCTTGTTCTTGGTTGGTCAGATGTTC
TCTTTYCAGCCTCGGCGCCACTGGACTACCCAGGATTGCAATTGTTCCATC



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Fig.3E

SEO ID NO. 43 (VN4, 7c) ATGAGCACACTTCCAAAACCCCAAAGAAAAACCAAAAGAAACACCATCCGCCGCCCACA GGACGTCAAGTTCCCGGGTGGCGGCCAGATCGTTGGTGGAGTCTACTTGCTGCCGCGCAG GGGCCCGCGCTTGGGTGTGCGCGCGACGAGAAAGACTTCTGAACGGTCCCAGCCCAGAGG TAGGCGCCAACCAATACCCAAAGTGCGCCACCAAACGGGCCGTACCTGGGCCCAGCCCGG CCGCGGCTCTCGCCCAAATTGGGGCCCAAACGACCCCGGGGAGGTCCGCAACTTGGG TAAAGTCATCGACACCCTTACTTGCGGCCTTCGCCGACCTCATGGGGTATATCCCTGTCGTAG GCGCTCCGWTGGGAGGCGTCGCGGNGGCCTTGGCGCATGGGGTCANGGNCATCGAGGACGGNGTAA ATTACGCAACAGNGAATCTTCCCGGNNGCTCTNTCTCTATCTTNCTCTTGGCACTTCTCTCGTGCC TTACAACACCAGCCTCCGCGGCGCATTATACCAACAAGTCTGGCCTGTACCATCTCACCAACGACT GCCCCAACAGCAGCATCGTTTATGAGGCGGAGACACTGATTTTGCACTTGCCTGGGTGTGTACCTT GTGTGAAGRTGRACAATCAATCCCGGTGCTGGGTGCAGGCCTCCCCGACCCTGGCAGTGCCGAACG CGTCTACGCCAGTCACCGGGTTCCGCAAACATGTGGACATCATGGTGGGCGCTGCCGCGTTCTGTT CAGCTATGTATGTGGGGGGACCTGTGCGGGGGCCTTTTCCTCGTTGGACAGCTCTTCACGCTCAGGC CTCGGATGCATCAGGTTGTCCAGGAGTGTAACTGTTCCATCTACACAGGGCATATCACTGGACACC GAATGGCA

SEO ID NO. 41 (FR1, 9a) ATGAGCACACTTCCAAAACCCCAAAGAAAAACCAAAAGAAATACTAACCGTCGCCCTATGGAC GTCAAGTTCCCGGGCGGCGGCCAGATCGTTGGTGGAGTTTACTTGTTGCCGCGCAGGGGC CCTCGTTTGGGTGTGCGCGCGACGAGAAGACCTCCGAACGGTCCCAGCCTAGAGGCAGG CGCCAGCCCATACCAAAGGTACGCCAGCCGACAGGCCGTAGCTGGGGTCAACCCGGCTAC CCTTGGCCCCTTTATGGCAACGAGGGCTGCGGATGGCCGGGATGGCTCCTGTCCCCCCGC GGGTCTCGTCCTAATTGGGGCCCCAACGACCCCCGGCGAAGGTCCCGCAACTTGGGTAAG GTCATCGATACCCTTACATNCGGNCTAGCCGACCTCATGGGGTACATCCCTGTCCTAGGAGG GCCGCTTGGCGGCGTTGCGGCTGCCCTGGCGCATGGCGTTAGGGCAATCGAGGACGGGGTCAATTA CGCAACAGGGAATCTTCCTGGTTGCTCCTTTTCTATCTTCCTCTTAGCACTGTTATCGTGCCTCAC TACACCAGCCTCAGCAATTCAAGTCAAGAACGCCTCTGGGATCTACCATCTTACCAATGACTGCTC GAACAACAGCATCGTTTTTGAGGCGGAGACCATGATACTGCATCTTCCAGGTTGTGTCCCATGTAT CAAGGCGGGGAATGAGTCACGATGTTGGCTCCCTGTCTCCCCCACCTTAGCCGTCCCCAACTCATC AGTGCCAATCCACGGGTTTCGCCGACACGTAGACCTCCTCGTTGGGGCAGCGGCATTTTGTTCGGC CATGTACATCGGAGACCTCTGTGGTAGCATAATCTTGGTAGGGCAGCTTTTTACTTTCAGGCCTAA GTACCATCAGGTTACCCAGGATTGTAACTGCTCTATNAACNCTGGCCACGTCACGGGACACAGGAT GGCA



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Fig.3F

SEQ ID NO. 49 (NE98, 10a)
ATGAGCACACTTCCTARACCACARAGARARACCARAGARACACCAACC?CCGGCCACAGGACGTT
AAGTTCCCAGGCGGCGGTCAGATCGTTGGTGGAGTTTACGTGCTACCACGCAGGGGCCCCCAGTTG
GGTGTGCGTGCAGTGCGCAAGACTTCCGAGCGGTCGCAACCCTGCAGTAGGCGCCCAACCCATCCCC
AGGGCGCCCCGAACCGAGGGCAGGTCCTGGGCTCAGCCCGGGTACCCTTGGCCCCTATATGGGAAT
GAGGCCTGCGGGTGGCAGGGTGGCTCCTGTCCCCGCGCGGTCCTC

SEQ ID NO. 53 (BNL1,1d)
CTCGACAGTTACTGAGAATGACATCCGTGTCGAGGAATCAATATACCAATGTTGTGACTTGGCCCC
CGAGGCTCGCAAGGCCATAAAGTCGCTCACCGAGCGGCTGTACATCGGGGGCCCYCTAACCAATTC
AAAAGGACAGAACTGCGGCTACCGTCGGTGCCGCGCGAGCGGCGTGCTGACTACCAGCTGCGGCAA
CACCCTGACATGCTACTTGAAAGCCAGAGCGGCCTGTCGAGCTGCAAAGCTCCGGGACTGCACCAT
GCTCGTGTGCGGGGATGACCTTGTCGTTATCTGTGAGAGTGCGGGAGTCGAGGAAGACGCGGCGAA
CCTACGAGCT

SEQ ID NO. 55 (BNL2,1d)
CTCGACAGTTACTGAGAACGACATCCGTACCGAGGRATCAATCTATCAATGTTGTGACTTGGCCCC
YGAGGCCCGCAAGGCCATAAAGTCGCTCACCGAGCGGTGTACGTCGGGGGCCCCCTAACCAATTC
AAAGGGGCAGAACTGCGGCTATCGTCGGTGTCGCGCTAGCGGCGTGCTGACCACCAGCTGCGGCAA
CACCCTCACATGCTACTTGAAAGCCAGGGCGGCCTGTCGAGCTGCAAAGCTCCAGGACTGCACGAT
GCTCGTGTGCGGAGACGACCTTGTCGTTATCTGTGAGAGCGCGGGAGTCGAGGAGGACGCGGCGAA
CCTACGAGTC

SEQ ID NO. 59 (CAM1078,1e)

CGTACAGCCTCCAGGACCCCCCTCCCGGAGAGCCATAGTGGTCTGCGGAACCGGTGAG
TACACCGGAATTGCCAGGACGACCGGGTCCTTTCTTGGATCAACCCGCTCAATGCCTGGA
GATTTGGGCGTGCCCCCGCAAGACTGCTAGCCGAGTAGTGTTTGGGTCGCGAAAGGCCTTG
TGGTACTGCCTGATAGGGTGCTTGCGAGTGCCCCGGGAGGTCTCGTAGACCGTGCACCAT
GAGCACCAATCCTAAACCTCAAAGAAAAACCAAAAGAAACACCAACCGCCGCCCACAGGA
CGTCAAGTTCCCGGGCGGTGGCCAGATCGTTGGTGGAGTCTACGTGCTACCGCGCAGGGC
CCCTAGATTGGGTGTGCGCAGCGCGGAAGACTTCGGAGGCCCGAACCTCGTGGGAG
CCCCTAGCTTATTCCCAAGGAGCGCCGACCCCAAGGTCCTGGCGCAGCCCGGGTA
CCCCTGGCCCCTCTATGGTAACGAGGGCTGCGGGTGGGCAGGTNGGCTCCTGTCCCCTCC
CGGCTCCCGTCCTAGTTGGGGTCTTACTGACCCCCGGCGTAGGTCACGCAATTTGGGTAA
GGTCATCGATACCCTCACGTGTTGNTTCGCCGACCTCATGGGGTACATACCG



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Fig.3G

SEQ ID NO. 61 (CAM1078, 1e)

CTCAACGGTCACTGAAGCTGATATCCGAACAGAGGAGTCCATATACCAATGCTGTGACCTGCACCC CGAAGCACGTGTAGCCATCAAGTCTTTGACTGAAAGGCTGTACGTCGGGGGGCCCTTGACCAATTC AAAAGGGGAGAACTGCGGCTATCGCAGATGCCGTGCCAGCGGCGTCTTGACAACCAGCTGCGGCAA CACCCTCACCTGCTATATCAAGGCCCTAGCAGCCTGTAGAGCTGCCAAGCTCCAGGACTGCACCAT GCTCGTCTGTGGCGACGACCTGGTCGTGATCTGCGAGAGTGTAGGGACCCAGGAGGATGCGGCGAG CCTGCGAGCC

SEQ ID NO. 63 (FR2, 1f)

NTCAACAGTCACTGAGAGTGATATCCGTACAGAGGAGTCCATCTACCAATGCTGTGATCTAGACCC CGAGGCTCGCAAGGCCATAAGGTCCCTCACAGAGAGGCTTTATATCGGGGGTCCCCTGACAAACTC AAAAGGGCAGAACTGCGCTACCGCCGATGCCGTGCAAGCGGCGTCCTGACGACTAGCTGCGGCAA CACCCTCACCTGTTACATAAAGGCCAGGGCAGCCTGTCGAGCTGCGAAGCTCCAGGATTGCTCAAT GCTCGTCTGTGGCGACGACCTTGTCGTTATCTGCGAGATCGAGGGGTCCANGAGGATCCGTCGAN NNNNNNNNNN

SEQ ID NO. 65 (FR16,1g)

SEQ ID NO. 67 (FR16,1g)

NNNNNNGTCACTGAGAGTGATATCCGTGTCGAGGARTCAATTTACCAATGCTGTGACCTGGCCCC CGAGGCTCGCGCGCGCACTCACCAACTC CGAGGCTCGCGCGAAAGTCGCTCACTGAGCGCTATATGTCGGGGGCCCTCTCACCAACTC AAAAGGACAGAACTGCGGCTATCGCCGGTGCCGTGCGAGCGGTGTGACTACTAGCTGCGGTAA CACCCTCACATGCTACCTGAAAGCCGCCGCGCCTGTCGAGCTGCAAAGCTCCGGGAATGCACAAT GCTCGTGTGTGGGGACGACCTCGTCGTTATCTGTGAGAGTGCGGGGGTCCAGGAGGATGCTGAAGCCTCGTNNNNNNN

SEQ ID NO. 69 (BNL3, 2e)

CTCGACAGTCACAGAGAGAGATATAAGNACTGAGGAGTCCATATACCAGGCTTGTTCCTTACCCGA GCAGGCCAGAACTGCCATACACTCATTGACTGAGAGACTCTACGTAGGAGGGCCCATGATGAACAG CAAAGGGCAATCCTGCGGATACAGGCATTGCCGCGCCAGCGGAGTGCTCACCACCAGTATGGGGAA TACCATCACGTGCTACATCAAGGCCCTAGCGGCTTGTAAAGCAGCAGGAATAGTGGCCCCCACCAT GCTGGTGTGCGGCGATGACCTAGTTGTCATCTCAGAGAGTCAGGGAGTCGAGGAGGACGACCGGAA CCTGANNNNN



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Fig.3H

SEQ ID NO. 71 (FR4, 2f)

CTCAACCGTCACAGAGAGGGATATAAGAACTGAGGAGTCCATATACCTGGCCTGCTCCTTACCCGA GCAGGCCCGGACTGCCATACATTCATTAACTGAGAGACTTTACGTGGGAGGGCCCATGATGAACAG CAAAGGGCAGTCCTGCGGATACAGGCGTTGCCGCGCTAGCGGAGTGCTCACCACCAGTATGGGGAA CACCATCACGTGTTATGTGAAAGCCCTCGCAGCTTGTAAAGCTGCGGGGCATTGTTGCCCCCACGAT GCTGGTGTGCGGCGATGACCTGGTTGTCATCTCAGAGAGTCAGGGGGGCTGAGGAGGACGAGCGAAA CCTGAGAGTC

SEQ ID NO. 73 (BNL5, 2h)

CTCAACAGTCGCGGAGAGAGACATCAGGACCGAGGAGTCCATTTACCTTGCCTGCTCCTTACCCGA GCAAGCCCGAACIGCCATACATTCATTGACTGAGAGACTTTACGTAGGAGGGCCCATGATGAACAG CAAGGGACAGTCCTGCGGTTACAGACGTTGCCGCGCGCGGGGGGGTGCTCACCACCAGCATGGGGAA TACCATCACATGCTATGTGAAGGCATTAGCTGCCTGCAAAGCTGCAGGCATCGTTGCTCCCACGAT GCTGGTTTGTGGCGACGATCTGGTCATCATCTCAGAGAGTCAGGGAACCGAGGAGGATGAGCGGAA CCTGAGAGTC

SEQ ID NO. 75 (FR13, 2k)

CGNACANCCTCCAGGCCCCCCCCCCCCGGGAGAGCCATAGTGGTCTGCGGAACCGGTGAG TACACCGGAATTGCCGGGAAGACTGGGTCCTTTCTTGGATAAACCCACTCTATGCCCGGC CATTTGGGCGTGCCCCGCAAGACTGCTARCCGAGTAGCGTTGGGTTGCGAAAGGCCTTG TGGTACTGCCTGATAGGGTGCTTGCGAGTGCCCCGGGAGGTCTCGTAGACCGTGCATCAT GAGCACAAATCCTAAACCTCAAAGAAAAACCAAAAGAAACACTAACCGCCGCCCACAGGA CGTTAAGTTCCCGGGCGGTGGCCAGATCGTTGGCGGAGTATACTTGTTGCCNTGCAGGGG NCCCAGGTNGNGTNTATGCGCAACGANGAAGACTNCCGAACAGTCCCAGCCACGTGGGAG GCGCCAGCCCATCCCGAAAGATCGGNGCACCACTGGCAAGTCCTGGGGACGTCCAGGATA TCCCTGGCCCTGTATGGGAACGAGGGCCTCGGGTGGCCAGGGTGGCTCCTGTCCCCCC GGGCTCCCGCCCGTCATGGGGCCCCACGGACCCCCGGCATAGGTCGCGCAACTTGGGTAA GGTCATCGATACCCTCACGTNCGGCTTTNCCGACCTCATGGGGTACATTCCCGTCGTTGG CGCCCAGTAGGNGGCGTCGCCAGAGCTCTCGCGCATGGCGTGAGAGTCCTGGAGGACGG TCTGTCCTGAATTACCGNGCCAGTTTCTGCTGTGGAAATCAAAAACACCAGMAACACATA CATGGTGACTAACGACTGTTCAAACAGYAGCATCACCTGGCAGCTTNNGNNCGCGGTGCT TCACGTTCCTGGATGCGTCCCCTGTGAACGAGGGGCAACAGTTCCCGGTGCTGGATTCC AGTCACGCCCRACGTAKNCGTGAGCCGACCTGGTGCCCTAACCGAGGGTTTGCGATCGCA CATCGACACCATCGTAGCGTCCGCAACATTTTGTTCTGCCCTCTACATAGGGGATGTATG TGGCGCGATAATGATAGCTGCCCAAGTGGTCATCGTCTCGCCGGAGCATCATCACTTTGT CCAGGACTGTAACTGTTCCATCTACCCGGGCCACATAACGGGGCCTCGTATGTNG

SEQ ID NO. 77 (FR13, 2k)

ATCCACAGTCACTGAAAGAGACATCAGAGTTGAAGAGTCCGTTTATCTGTCCTGTTCACTTCCCGA GGAGGCCCGAGCTGCCATACACTCACTAACTGAGAGGCTGTACGTGGGAGGTCCCATGCAGAACAG CAAGGGGCAATCCTGCGGATACAGGCGCTGCCGCGCGAGGGGGTGCTCACCACTAGCATGGGGAA TACTCTCACATGCTACTTGAAGGCCCAGGCGGCCTGCAGGGCCGCGGGCATTGTTGCACCCACAAT GCTGGTGTGTGGCGACGACCTGGTCGTCATCTCAGAGAGTCAGGGGACTGAGAGGGACGAGAACAA CCTGAGACCT



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Fig.3I

SEO ID NO. 79 (FR18,21)

CTCAACAGTCACGGAGAGGGACATCAGGAATGAGGAGTCCATATTCCTGGCCTGCTCGTTGCCCGA GGAGGCCCGGACTGTCATACATTCGCTCACTGAGAGACTCTACATAGGCGGGCCGATGATGAACAG CAAAGGCCAGTCCTGTGGATACAGGCGTTGTCGCGCCAGCGGGGTGTTCACCACTAGCATGGGCAA TACCATCACGTGCTATGTGAAAGCCATGGCAGCTTGCAGAGCTGCCGGGATTGACGCCCCCACAAT GTTGGTATGTGGCGACGACCTGGTGGTCATCTCAGAGAGTCAGGGGACCGAGGAGGACGAGCGAAA TCTGAGAGTC

SEQ ID NO. 81 (PAK64,3g)

CTCTTGACTCTACTGTCACTGAACAGGATATCAGGGTAGAAGAAGAAATATACCAATGTTGTGACC TTGAGCCGGAGGCTAGACGGCAATCAAATCGCTCACGGAACGGCTTTACGTTGGAGGTCCCATGT TCAACAGCAAGGGGCTCAAATGCGGATATCGCCGTTGCCGTGCTAGCGGTGTATTGCCCACTAGCT ACGCTAATACAATCACCTGCTACATCAAGGCCAGAGCGGCTGCTCGAGCTGCGGGCCTTCAAGACC CATCATTCCTTGTCTGCGGAGATGATTTGGTGGTAGTGGCTGAGAGTTGCGKCGTTGATGAGGAGG ATAGGGCAGC

SEO ID NO. 83 (BNL8, 4k)

CTCCACTGTAACCGAAAAGGACATCAGGCCCGAGGAAGAGGTCTATCAGTGTTGTGACCTGGAGCC CGAAGCTCGCAAGGTTATTACCGCCCTCACAGAAAGACTCTACGTGGGCGGCCCCATGCACAACAG CAAGGGAGACCTTTGTGGGTATCGGAGATGCCGCGCAAGCGGCGTCTACACGACCAGCTTCGGAAA CACACTGACGTGCTACCTCAAAGCCTCAGCTGCTATTAGAGCGGCAGGGCTGAGAGACTGCACCAT GCTGGTTTGCGGTGACGACTTGGTCGTCATCGCTGAGAGCGATGGCGTAGAGGAGGATAACCGAGC CCTCCNAGCC

SEQ ID NO. 85 (BNL12,41)

CTCCACGGTGACTGAAAAGGACATCAGGGTCGAGGAAGAGATCTATCAATGTTGTGACCTGGARCC CGAAGCCCGCAAAGCAATATCCGCCCTCACAGAGAGRCTCTACTTGGGCGGCCCCATGTATAACAG CAAAGGGGAGCTCTGCGGGTATCGGAGGTGCCGCGCGAGCGGAGTGTACACCACAAGTTTCGGGAA CACRETGACCTGCTATCTTAAGGCCACCGCAGCTACCAGGGCTGCAGGCCTAAAAGACTGCACCAT GCTGCTCTGCGGTGACGACTTGGTCGTCATCGCCGAGAGCGAGGGCGTAGAGGAGGATTCCCAACC CCTCCGAGCC

SEQ ID NO. 87 (EGE1, 4m)

CTCCACCGTAACCGAAAGGGACATCAGGGTCGAGGAGGAGGTCTATCAGTGTTGTGATCTGGAGCC AGAGGCCCGCAAGGCAATATCCGCCCTCACGGAGAGACTCTATGTGGGCGGTCCCATGTTTAACAG CAAGGGAGACCTATGTGGCTACCGCAGGTGCCGCGCAAGCGGCGTCTACACCACCAGCTTCGGAAA GCTGGTTTGCGGGGACGACCTGGTCGTCATCGCAGAGAGCGATGGCGTGGACGAGGACCGCCGAGC CCTCCAAGCT

SEO ID NO. 89 (VN13,7a)

CTCAACAGTCACAGAGCGCGATGTCCAGACGGAGCATGACATCTACCAGTGCTGTAAGTTGGAGCC CGCAGCACGGACAGCCATCACATCGCTTACTGACCGATTGTACTNCGGTGGTCCCATGTNTAACTC TAARGGTCAGGCATGTGGATACCGTAGGTGCAGGGCCAGTGGCGTCTTGACCACCATCCTGGCCAR TACTCTGACTTGCTACTTGAAAGCTCAGGCGGCATGCAGAGCTGCCGGGCTGAAGGACTTTGACAT GTTGGTCTGCGGAGACGACCTTGTCGTTATTTCGGAGAGTTTGGGGGTCTCGGAGGACACTAGTGC ACTGCGAGCT



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Fig.3J

SEQ ID NO. 91 (VN4,7c)

CTCGACAGTCACCGAGCGCGACATCCRCACCGAGCACGACATCTACCAATGCTGCCAACTTGACCC
GGTGGCACGCAAGGCTATTACATCTCTGACTGAGCGGCTGTACTGCGGWGGGCCCATGATGAACTC
CCGTGGTCAATCATGTGGATACCGTAGGTGCCGAGCCAGTGGCGTGCTCACCACGAGCTTGGGCAA
TACCCTAACATGCTATTTGAAAGCACAAGCAGCGTGTAGGGCAGCAAAGCTCAAAAACTATGACAT
GTTAGTCTGCGGAGACGATCTAGTCGTTATCGCGGAGAGTGGAGGAGTCTCTGAGGATGTTGACGC
CCTGCCGAGCA

SEQ ID NO. 93 (VN12,7d)

CTCCTCCGTCACGGAGCGTGACATCCGCACTGAACACGACATCTATCAGTGCTGCCAATTAGATCC
GGTAGCACGGAAAGCCATTACATCTCTTACTGAGCGGCTGTACTGCGGCGCCCCCATGTACAACTC
TCGAGGTCAGTCATGTGGGTACCGCAGGTGCCGGGCTAGTGGTGTCTTCACCACAAGCTTGGGCAA
CACCATGACATGCTACCTGAAGGCTCAGGCGGCTTGTAGGGCAGGCRAAGCTCAAAAACTTTGACAT
GTTGGTCTGCGGAGACGACCTAGTCGTTATTGCTGAGAGCGGAGGAGTCCCTGAGGATGCCGGGGC
CCTGCGAGTC

SEQ ID NO. 95 (FR1,9a)

ATCCACAGTCACGGGGCGCGACATACGCACAGAACNAGACATTTACCTGTCCTGCCAGCTCGACCC AGAGGCCCGGAAAGCCATAAAGTCTCTCACTGAGAGGCTCTATGTCGGGGGCCCTATGTACAACTC AAAGGGCCAACTCTGTGGTCAACGCCGATGCCGAGCAAGCGGAGTACTCCCCACAAGCATGGGTAA CACCATCACATGCTTCCTGAAGGCAACCGCCGCTTTGCCGAGCAGCCGGCTTTACAGATTATGACAT GTTGGTCTGCGGAGACGATTTGGTTGTCGTAACTGAGAGTGCTGGAGTCAACGAGGATATCGCTAA CCTGCGAGCC

SEQ ID NO. 97 (NE98, 10a)

SEQ ID NO. 99 (FR14,11a)

SEQ ID NO. 101 (FR15,11a)



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Fig.3K

SEQ ID NO. 103 (FR19,11a)
CGTACAGCCTCCAGGACCCCCCCTCCCGGGAGGCCATAGTGGTCTGCGGAACCGGTGAGTACACC
GGAATTGCCGGGAAGACTGGGTCCTTTCTTGGATTAACCCACTCTATGCCCGGAGATTTGGGCGTG
CCCCCGCAAGACTGCTAGCCGAGGTAGCGTTGGGTTGCGAAAGGCCTTGTGGTACTGCCTGAAACC
TGCTTGCGAGTGCCCCGGGAGGTCTCGTAGACCGTGCACCATGAGCACCAATCCTAAACC
ACAAACCAAAAGAAACACCAACCGCCGCCCCACAGGACGTTTAGTTCCCGGGCGGTGGCCAGATCGT
TGGCGGGGTGTACTTGTTGCCGCGCAGGGGCCCCAACCTATCCCCAAGGTTAGGCGCACCACCACCGGCCGTT
GGAGCGGTCCCAGCCGCGTGGGAGGCGCCCAACCTATCCCCAAGGTTAGGCGCACCACCACCGGCCGTT

SEQ ID NO. 105 (FR19,11a)
CTCTACTGTCACAGAGAGGGATATACGAACAGAGGAATCCATYTATCTGGCTTGTCAATTGCCCGA
AGAGGCCCGGAAGGCCATCAAATCACTGACAGAGAGACTATACGTGGGCGCCCGATGGAAAACAG
CARGGGCCAGGCCTGCGGATACAGGCGTTGCCGCGCAAGCGGGGTATTCACCACAAGCTTGGGGAA
CACCATGACTTGTTACATCAAAGCCAAGGCGGCTTGTAAAGCCGCTGGCATTGTTGACCCAGTGAT
GCTCGTGTGCGGCGACGACCTAGTGGTCATCTCAGAAAGCAAGGGGGGTGGAGGAGGACCAACGAGA
CCTACGANTC

SEQ ID NC. 2 (BNL1, 1d)
MSTNPKPQRKTKRNTNRRPXXXXXPGGGQIVGGVYLLPRRGPRXGVRATRKTSERSQPRGRRQPIP
KAXRXEGRSWAQPGYPWPLYGNEGCGWAXWLLSPRGSRPNWGP

SEQ ID NO. 4 (BNL1, 1d)
DGVNYATGNLPGCSFSIFLLALLSCLTVPXTAHEVRNASGVYHVTNDCSNSSIIYEMDGMIMHYPG
CVPCVREDNHLRCWMALTPTLAVKXASVPTXAIRRHVDLLVGXXTFCSAMYVXDLCGSVFLAGQLF
TFSPRMHHTTQECNCSI

SEQ ID NO. 6 (BNL2, 1d)
MSTNPKPQRKTKRNTNRRPQDVKXPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRDRRQPIP
KARQSDGXXWAQPGHPWPLYGNEGCGWAGWLLSPRGSRPSWGP

SEQ ID NO. 8 (BNL2, 1d)
DGVNYATGNLPGCSFSIFLLAFLSCLTVPTTAHEVRNASGVYHLTNDCSNSSIIYEMSGMILHAPG
CVPCVRENNSSRCWMXLTPTLAVKDANVPTAAIRRHVDLLVGTAAFRSAMYVGDLCGSVFLVGQLF
TFSPRLYHTTGECNCSI

SEQ ID NO. 10 (CAM1078, 1e)
MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYVLPRRGPRLGVRAARKTSERSQPRGRRQPIP
KERRPEGR

SEQ ID NO. 12 (FR2, 1f)
MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP
KARRPEGRSWAQPGYPWPLYANEGCGWAGWLLSPRGSRPSWGPNDPRRSRNLGKVIDTLTCGFAD
LMGYIPLVGAPLGGASRTLXHGVRVLXGGVXXXXXNLXGCSXXIFLLXLLSCLTVPTSAYEVHSTT
DGYHVTNDCSNGSIVYEAKDIILHTPGXVPCIREGNISRCWVPLTPTLAARIANAPIDEVRRHVDL
LVGAAVTCSAMYIGDLCGGVFLVGQLFTFTSRRHWT
VODCNCSIYSGHITGHXXX

SEQ ID NO. 14 (BNL3, 2e)
MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP
KDRXATGRSWGRPGYPWPLYGNEGLGWAGWLLSPRGSRPSWG

SEQ ID NO. 16 (BNL3, 2e)
TCXXADLMGYXPVVGAPVGGXARALAXGVRVLEDGINYXTGNLPGCSFSIFXLALLSCVTVPVSXV
TCXXADLMGYXPVVGAPVGGXARALAXGVRVLEDGINYXTGNLPGCSFSIFXLALLSCVTVPVSXV
EVKNTSQAYMATNDCSNNSIVWQLXDAVLHVPGCVPCENSSGRFHCWIPISPNIAVSKPGALTKGL
RARIDAVVMSATLCSALYVGDVCGAVMIAAQAFIVAPKRHYFVQECNCSIYPGHITGHRMA



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Fig.3L

SEQ ID NO. 18 (FR4, 2f)
MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRAPRKTSERSQPRGRRQPIP
KDRRATGKSWGRPGYPWPLYGNEGLGWAGWLLSPRGSRPSWGPNDPRHRSRNLGKVIDTLTCGFXD
LMGYIPVVGAPVGGVARALAHGVRVLEDGINYATGNLPGCSFSIFLLALLSCITVPVSAIQVKNNS
HFYMATNDCANDSIVWQLRDAVLHVPGCVPCERSGNRTFCWTAVSPNVAVSRPGALTRGLRAHIDT
IVMSATLCSALYIGDLCGAVMIAAQVAVVSPQYHTFVQECNCSIYPGHITGHRMX

SEQ ID NO. 20 (BNL4, 2g)
DGVNYATGNLPGCSFSIFLLALLSCVTVPVSAVQVKNTSTMYMATNDCSNNSIIWQMQGAVLHVPG
CVPCELQGNKSRCWIPVTPNVAVNQPGALTRGLRTHIDTIVMVATLCSALYIGDVCGAVMIAAQVV
IVSPQHHNFSQDCNCSI

SEQ ID NO. 22 (BNL5, 2h)
MSTNPKPQRKTKRNTNRRPQDVKFPGGGRSLAEYTCARRGKLRRSSMG

SEQ ID NO. 24 (BNL5, 2h)
DGINYATGNLPGCSFSIFLLALLSCLTVPASAVQVKNTSHSYMVTNDCSNSSIVWQLKDAVLHVPG
CVPCERHQNQSRCWIPVTPNVAVSQPGALTRGLRTHIDTIVASATVCSALYVGDFCGAVMLVSQFF
MISPQHHIFVQDCNCSI

SEQ ID NO. 26 (BNL6, 21)
DGINYATGNLPGCSFSIFLLALLSCITVPVSAVQVANRSGSYMVTNDCSNSSIVWQLEEAVLHVPG
CVPCEWKDNTSRCWIPVTPNIAVSQPGAXTKGLRTHIDIIVASATFCSALYV

SEQ ID NO. 28 (BNL7, 4k)
MSTNPKPQRKTKRNTNRRPMDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP
KARRSEGRSWAQPGYPWPLYGNEGCGWAXWLLSPRGSRPSWGPNDPRRSR

SEQ ID NO. 30 (BNL7, 4k)
DGINFATGNLPGCSFSIFLLALLSCLTVPASAINYRNVSGIYYVTNDCPNSSIVYEADHHILHLPG
CVPCVREGNÇSRCWVALTPTVAAPYIGAPLESLRSHVDLMVGAATVCSALYIGDXCXGLFLVGQMF
SFRPRRHWTTQDCNCSI

SEQ ID NO. 32 (BNL8, 4k)
DGINYATGNI PGCSFSIFILALLSCLTVPASAINYRNTSGIYHVTNDCPNSSIVYEADHHILHLPG
CVPCVRTGNQSRCWVALTPTVAAPYIGAPLESLRSHVDLMVGAATVCSALYIGDLCGGUFLVGQMF
SFRPRRHWTAQDCNCSI

SEQ ID NO. 34 (BNL9, 4k)
DGINYATGNLPGCSFSIFLLALLSCLTVPASAINYHNTSGIYHITNDCPNSSIVYEADHHILHLPG
CVPCVRVGNQSSCWVALTPTIAAPYIGAPLESIRSHVDLMVGAATVCSALYIGDLCGGAFLVGQMF
SFRPRRHWTTQDCNCSI

SEC ID NO. 36 (BNL10, 4k)
DGINYATGNIPGCXFSIFLXALLSCLTVPASATNYRNVSGIYHVTNDCPNSSIVYEADHHILALPG
CVPCVRVGNQSRCWVALTPTVAAPYTAAPLESLRSHVDLMVGAATVCSALYIGXLCGGLFLVGQMF
SXQPRRHWTTQDCNCSI

SEQ ID NO. 38 (BNL11, 4k)
DGINYATGXLPGCSFSIFLLALLSCLTVPASATNYRNVSGIYHVTNDCPNSSIVFEADHHILHLPG
DGINYATGXLPGCSFSIFLLALLSCLTVPASATNYRNVSGIYHVTNDCPNSSIVFEADHHILHLPG
CVPCVKEGNHSRCWVALTPTVAAPYIGAPLESLRSHVDVMVGAATVCSALYIGDLCGGLFLVGQMF
SFRPRRHWTTQECNCSI

SEQ ID NO. 48 (BNL12, 41)
DGINYATGNLPGCSFSIFILALLSCLTVPASAQHYRNVSGIYHVTNDCPNSSIVYESDHHILHLPG
CVPCVKTGNTSRCWVALTPTVAAPILSAPLMSVRRHVDLMVGAATLSSALYVGDLCGGAFLVGQMF
TFQPRRHWTVQDCNCSI



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Fig.3M

SEQ ID NO. 46 (VN13, 7a)

MSTLPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP KVRHQTGRTWAQPGYPWPLYGNEGCGWAGWLLSPXGSRPNWGPNDPRXRSRNLGKVIDTLTXXFAD LIEYI

SEQ ID NO. 44 (VN4, 7c)

MSTLPKPQRKTKRNTIRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP KVRHQTGRTWAQPGYPWPLYGNEGCGWAGWLLSPRGSRPNWGPNDPRRRSRNLGKVIDTLTCGFAD LMGYIPVVGAPXGGVAXALAHGVXXIEDXVNYATXNLPXXSXSIXLLALLSCLTTPASAAHYTNKS GLYHLTNDCPNSSIVYEAETLILHLPGCVPCVKXXNQSRCWVQASPTLAVPNASTPVTGFRKHVDI MVGAAAFCSAMYVGDLCGGLFLVGQLFTLRFRMHQVVQECNCSIYTGHITGHRMA

SEQ ID NO. 48 (VN12, 7d)

MSTLPKPQRKTKRNTNRRPMDVKFPGGGQIVGGVYLLFRRGPRLGVRATRKTSERSQARGRRQPIP KVRQNQGRTWAQPGYPWPLYGNEGCGWAGWLLSPRGSRPDWXPNDPRXRSRNLGKVIDTLTCGFAD LMEYIPVVGAPLGGVAAELXHGVRAIEDGINYATGNLPGCSFSIFXLALLSCLTTPASALNYANKS GLYHLTNDCFNSSIVYEANGMILHLPGCVPCVKTGNLTKCWLSASPTLAVQNASVSIRGVREHVDL LVGAAAFCSAMYVGDLCGGLFLVGQLFTFRPRMYEIAQDCNCSIYAGHITGHRMA

SEQ ID NO. 42 (FR1, 9a)

MSTLPKPQRKTKRNTNRRPMDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP KVRQPTGRSWGQPGYPWPLYGNEGCGWAGWLLSPRGSRPNWGPNDPRRRSRNLGKVIDTLTXXLAD LMGYIPVLGGPLGGVAAALAHGVRAIEDGVNYATGNLPGCSFSIFLLALLSCLTTPASAIQVKNAS GIYHLTNDCSNNSIVFEAETMILHLPGCVPCIKAGNESRCWLPVSPTLAVPNSSVPIHGFRRHVDL LVGAAAFCSAMYIGDLCGSIILVGQLFTFRPKYHQVTQDCNCSXNXGHVTGHRMA

SEQ ID NO. 50 (NE98, 10a)

MSTLPKPQRKTKRNTNXRPQDVKFPGGGQIVGGVYVLPRRGPQLGVRAVRKTSERSQPRSRRQPIPRARRTEGRSWAQPGYPWPLYGNEGCGWAGWLLSPRGSRPSWGPNDPRRR

SEQ ID NO. 52 (NE98, 10a)

DGINFATGNLPGCSFSIFLLALFSCLLTPTAGLEYRNASGLYMVTNDCSNGSIVYEAGDIILHLPG CVPCVRSGNTSRCWIPVSXTVAVKSPCAATASLRTHVDMMVXAATLCSALYVGDLCGALFLXGQGF SWRHROHWTVODCNCSI

SEQ ID NO. 54 (BNL1,1d)

STVTENDIRVEESIYQCCDLAPEARKAIKSLTERIYIGGXLTNSKGQNCGYRRCRASGVLTTSCGN TLTCYLKARAACRAAKLRDCTMLVCGDDLVVICESAGVEEDAANLRA

SEQ ID NO. 56 (BNL2, ld)

STVTENDIRTEXSIYQCCDLAXEARKAIKSLTERLYVGGPLTNSKGQNCGYRRCRASGVLTTSCGN TLTCYLKARAACRAAKLQDCTMLVCGDDLVVICESAGVEEDAANLRV

SEO ID NO. 58 (FR17,1d)

STVTENDIRVEESIYQCCDLAPEARKAIKSLTERLYIGGPLTNSKGQNCGYRRCRASGVLTTSCGN TLTCYLKARAACRAAKLQDCTMLVCGDDLVVICESYGVEEDAANLRV



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Fig.3N

SEO ID NO. 60 (CAM1078, 1e)

MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYVLPRRGPRLGVRAARKTSERSQPRGRRQPIP KERRPEGRSWAQPGYPWPLYGNEGCGWAGXLLSPRGSRPSWGPTDPRRRSRNLGKVIDTLTCXFAD LMGYIP

SEQ ID NO. 62 (CAM1078,1e)

STVTEADIRTEESIYQCCDLHPEARVAIKSLTERLYVGGPLTNSKGENCGYRRCRASGVLTTSCGN TLTCYIKALAACRAAKLQDCTMLVCGDDLVVICESVGTQEDAASLRA

SEQ ID NO. 64 (FR2, 1f)

STVTESDIRTEESIYQCCDLDPEARKAIRSLTERLYIGGPLTNSKGQNCGYRRCRASGVLTTSCGN TLTCYIKARAACRAAKLQDCSMLVCGDDLVVICEIEGXXEDPSXXXX

SEQ ID NO. 66 (FR16,1g)

MSTNPKPQRKTKRNINRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSERSQPRGRRQPIP KARRSEGRSWAQPGYPWPLYGNEGMGWAGWLLSPHGSRPSWGPSDPRRRSRNLGKVIDTLTCGFAD LMGYIPLVGAPLGGVARALAQGFRDL

SEQ ID NO. 68 (FR16,1g)

XXVTESDIRVEXSIYQCCDLAPEARVAIKSLTERLYVGGPLTNSKGQNCGYRRCRASGVLTTSCGN TLTCYLKAAAACRAAKLRECTMLVCGDDLVVICESAGVQEDAASXXX

SEQ ID NO. 70 (BNL3, 2e)

STVTERDIXTEESIYQACSLPEQARTAIHSLTERLYVGGPMMNSKGQSCGYRHCRASGVLTTSMGN TITCYIKALAACKAAGIVAPTMLVCGDDLVVISESQGVEEDDRNLXX

SEQ ID NO. 72 (FR4, 2f)

STVTERDIRTEESIYLACSLPEQARTAIHSLTERLYVGGPMMNSKGQSCGYRRCRASGVLTTSMGN TITCYVKALAACKAAGIVAPTMLVCGDDLVVISESQGAEEDERNLRV

SEQ ID NO. 74 (BNL5, 2h)

STVAERDIRTEESIYLACSLPEQARTAIHSLTERLYVGGPMMNSKGQSCGYRRCRASGVLTTSMGN TITCYVKALAACKAAGIVAPTMLVCGDDLVIISESQGTEEDERNLRV

SEO ID NO. 76 (FR13,2k)

MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLXCRXPRXXXCATXKTXEQSQPRGRRQPIP KDRXTTGKSWGRPGYPWPLYGNEGLGWACWLLSPRGSRPSWGPTDPRHRSRNLGKVIDTLTXGFXD LMGYIPVVGAPVXGVARALAHGVRVLEDGINYETGNLPGCSFSISLLALLSITXPVSAVEIKNTXN TYMVTNDCSNXSITWQLXXAVLHVPGCVPCEREGNSSRCWIPVTPXVXVSRPGALTEGLRSHIDTI VASATFCSALYIGDVCGAIMIAAQVVIVSPEHHHFVQDCNCSIYPGHITGPRMX

SEO ID NO. 78 (FR13,2k)

STVTERDIRVEESVYLSCSLPEEARAAIHSLTERLYVGGPMQNSKGQSCGYRRCRASGVLTTSMGN TLTCYLKAQAACRAAGIVAPTMLVCGDDLVVISESQGTERDENNLRP



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Fig.30

SEQ ID NO. 80 (FR18, 21)

STVTERDIRNEESIFLACSLPEEARTVIHSLTERLYIGGPMMNSKGQSCGYRRCRASGVFTTSMGN TITCYVKAMAACRAAGIDAPTMLVCGDDLVVISESQGTEEDFRNLRV

SEQ ID NO. 82 (PAK64,3g)

STVTEQDIRVEEEIYQCCDLEPEARRAIKSLTERLYVGGPMFNSKGLKCGYRRCRASGVLPTSYGN TITCYIKARAARRAGLQDPSFLVCGDDLVVVAESCXVDEEDRAALR

SEO ID NO. 84 (BNL8,4k)

STVTEKDIRPEEEVYQCCDLEPEARKVITALTERLYVGGPMHNSKGDLCGYRRCRASGVYTTSFGN TLTCYLKASAAIRAAGLRDCTMLVCGDDLVVIAESDGVEEDNRALXA

SEQ ID NO. 86 (BNL12,41)

STVTEKDIRVEEEIYQCCDLXPEARKAISALTEXLYLGGPMYNSKGELCGYRRCRASGVYTTSFGN TVTCYLKATAATRAAGLKDCTMLVCGDDLVVIAESEGVEEDSQPLRA

SEQ ID NO. 88 (EG81, 4m)

STVTERDIRVEEEVYQCCDLEPEARKAISALTERLYVGGPMFNSKGDLCGYRRCRASGVYTTSFGN TLTCYLKATAATRAAGLKDCTMLVCGDDLVVIAESDGVDEDRRALQA

SEQ ID NO. 90 (VN13,7a)

STVTERDVQTEHDIYQCCKLEPAARTAITSLTDRLYXGGPMXNSKGQACGYRRCRASGVLTTILAN TLTCYLKAQAACRAAGLKDFDMLVCGDDLVVISESLGVSEDTSALRA

SEQ ID NO. 92 (VN4,7c)

STVTERDIXTEHDIYQCCQLDPVARKAITSLTERLYCXGPMMNSRGQSCGYRRCRASGVLTTSLGN TLTCYLKAQAACRAAKLKNYDMLVCGDDLVVIAESGGVSEDVDALRA

SEQ ID NO. 94 (VN12,7d)

SSVTERDIRTEHDIYQCCQLDPVARKAITSLTERLYCGGPMYNSRGQSCGYRRCRASGVFTTSLGN TMTCYLKAQAACRAXKLKNFDMLVCGDDLVVIAESGGVPEDAGALRV

SEQ ID NO. 96 (FR1,9a)

STVTGRDIRTEXDIYLSCQLDPEARKAIKSLTERLYVGGPMYNSKGQLCGQRRCRASGVLPTSMGN TITCFLKATAACRAAGFTDYDMLVCGDDLVVVTESAGVNEDIANLRA

SEQ ID NO. 98 (NE98, 10a)

STVTEQDIRVELSIFQACDLKDEARRVITSLTERLYCGGPMFNSKGQHCGYRRCRASGVLPTSFGN TITCYIKAKAATKAAGIKNPSFLVCGDDLVVIAESAGIDEDKSALRA

SEQ ID NO. 100 (FR14,11a)

STVTERDIRTEESIYLSCQLPEEARKAIKSLTERLYVGGPMENSKGQACGYRRCRASGVFTTSLGN TMTCYIKAKAACKAAGIVDPVMLVCGDDLVVISESKGVEEDQRDLRV



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Fig.3P

Figure 3 - continued

SEC ID NO. 102 (FR15,11a)

STVTERDIRTEESIXXACQLPEEARKAIKSLTERLYVGGPMENSKGQACGYRRCRASGVFTTSLGN TMTCYIKAXAACKXAGIVDPVMLVCGDDLVVISESKGVEEDQRDLXX

SEQ ID NO. 104 (FR19,11a)

 ${\tt MSTNPKPQRQTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRVGVRATRKTSERSQPRGRRQPIPKVRRTTGR}$

SEQ ID NO. 106 (FR19,11a)

STVTERDIRTEESXYLACQLPEEARKAIKSLTERLYVGGPMENSKGQACGYRRCRASGVFTTSLGN TMTCYIKAKAACKAAGIVDPVMLVCGDDLVVISESKGVEEDQRDLRX



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FIG. 4A	Core/El		amino acid alignment
Isolate	Type	SEQ	
HCV-1	<u>-</u>	229	L ACTIVIDADE DE L'AUTOPOS DE L'ACTIVIS DE L'
HCV-J	1p	230	
BNL1	1q	2	XXXXXX
BNL2	19	9	XX
CAM1078	1e	10/60	R-T
FR2	1 £	12	
FR16	19	99	R-TI
HC-J6	2a	231	
HC-J8	5 p	232	
CH610	2c	3	
NE92	2d	234	
BNL3	5е	14	
FR4	2£	18	R-TP-
FR13	2k	9/	R-TXXXCX
EB1	3a	247	R-TI
NZL1	3a	248	LR-TI
HCV-TR	3p	235	LRQTLN
GB358	4°C	249	
DK13	4 d	236	R-TM
CAM600	4e	237	R-TM
GB809	4 e	238	L-R-T
HPCCOREEZA	4.2	250	5
HPCCOREZB	45	251	W
HPCCOREZC	43	252	
GB724	4.5	253	R-TMM
BNL7	4 4 4	28	R-TM
BE95	5a	239	R-T
HK2	6a	240	LR-TT
VN13	7a	46	LR-T
VN4	7c	44	LR-TI
VN12	7d	48	LR-TM
FR1	9a	42	LR-T
NE98	10a	20	LR-TXV-
FR19	11a	104	RQTV

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4B
Fig.

Isolate	Туре	SEQ	Core-V
			51 100
HCV-1	٦a	229	KTSERSQPRGRRQPIPKARRPEGRTWAQPGYPWPLYGNEGCGWAGWLLSP
HCV-J	1p	230	
BNL1	1q	7	XXSX-X-X
BNL2	1q	9	
CAM1078]e	10/60	(A)
FR2	1 £	12	
FR16	19	99	
HCJ6	2a	231	
HCJ8	2 p	232	
CH610	2c	233	
NE92	2d	234	T-RS-GKT-L-KS-GK
BNL3	2e	14	
FR4	2£	18	DAT-KS-GR
FR13	2k	9/	X-QLD-XTT-KS-GRL
EB1	3а	247	SSS
NZL1	3а	248	
HCV-TR	3p	235	KO-HLSRSKO-HL
GB358	4c	249	
DK13	4 d	236	
CAM600	4e	237	
GB809	4 e	238	
BNL7	4 k	28	X
HPCCOREEZA	4 ?	250	SSF-
HPCCOREZB	4?	251	
HPCCOREZC	4?	252	XSS
GB724	4.2	253	
BE95	5 a	239	
HK2	6a	240	
VN13	7a	46	
VN4	7c	44	V-HQT
VN12	7d	48	
FR1	9a	42	
NE98	10a	20	SSRTS
0100	112	70,	11 mm



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Isolate	Type		101
HCV1	la	229	rgsrpswgptdprrrsrnlgkvidtltcgfadlmgyiplvgaplggaara
HCV-J	1b	230	
BNL1	1d	~	N
BNL2	1d	9	
CAM1078	1e	10/6	10/60X
FR2	1£	12	L-SNN
FR16	19	99	Λ
HC-J6	2a	231	^
HC-J8	Sp	232	T
CH610	2c	233	· \-\-\-\-\-\\
NE92	2d	234	\-\\
BNL3	2e	14	SEQ ID NO: 16XXX-VX-V-
FR4	2£	18	\n\n\n\n\n\n\n-
FR13	2k	9/	\X\\XX
HCV-TR	3р	235	
GB116	4°C	241	
DK13	4 d	236	\ \ \ \ \ \ \ \ \ \ \ \ \
CAM600	4 e	237	
GB809	4 e	238	1 1 1 1 1 1
G22	4 £	242	· · · \ \ - \ \ - \ \ - · · · · · · · ·
GB549	49	243	\ \
GB438	4 h	244	'
BNL7	4 k	28	
BE95	Sa	239	
HK2	6а	240	HN
VN13	7a	46	XX
VN4	7c	44	-X-NXN
VN12	7d	48	D-X-NX
FR1	9a	42	NNXXLXXLVL-GV-A-
NE98	10a	50	



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9	
7.	
Fig	

Isolate

3	77.) [141
		1	151
HCV1	19	229	GVRVLEDGVNYATGNLPGCSFSTF1.1.A1.1.SCT.TVPASAVOVRNSTG
HCV-J	1p	230	
BNL1	1d	4	AS
BNL2	1d	œ	- F
FR2	1£	12	-XXGXXXXXXXXXTE-HST-DG
FR16	19	99	Q-F-D-
HC-J6	2a	231	
HC-J8	5 p	232	IISSS
CH610	2c	233	III
S83	2c	254	VE - KDTGDS
NE92	2d	234	II
BNL3	2e	16	XIXXXV-XVE-K-TSQA
FR4	2£	18	HSN-XIVIK-NSHF
BNL4	<u>2</u> g	20	MLST-XVVVK-TSTM
BNL5	2h	24	SHST-X
BNL6	2i	56	IVVVRS-S
FR13	2k	97	I-X-VVEIK-TXNT
BR36	3а	255	LEWTS
HCV-TR	3p	235	A-GTSFGLEYT-TS
24	4a	256	EHYAS-I
GB809-4	4a	257	EHYAS-I
21	4 b	258	VHYAS-V
GB116	4 _C	241	-EAVI
GB215	4°C	259	IHYAS-V
GB358	4c	260	VNY AS-I
DK13	4q	236	I
CAM600	4 e	237	AVIIIAS-I
GB809-2	4 e	238	AVIAS-V
CAMG22	4 £	261	AVI
CAMG27	4 £	262	I-ST-HYHV
GB549	49	243	AVI
GB438	4 h	244	AVIQHYAS-I
BNL7	4 k	30	I-FINYVS-I
BNL8	4 4	32	I-STXII
BNL9	4 k	34	I-SL-HANII
BNI.10	4 k	36	1-SVVNTXXTT



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BNL11	4 k	4k 38	I-SAXNLXXI
BNL12	41	40	III
BE95	5a	239	
BE100	5a	263	
HK2	6a	240	
VN4	7c	44	XXIXXIXXXX-X-XXAHYT-KS
VN12	7d	48	-XAIIINYA-KS
FR1	9a	42	I-SK-X
NEGO	0	C	0



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			Fig. 4F	
Isolate	Туре	SEQ	V1 V3 V4	
HCV-1	1a	229	TNDCPNSSIVYEAADAILHTPGCVPCVREGNASRCWVAMTPTVATR	
HCV-J	1p	230	W-W	
BNL1	ָרָק די	4r (H-QSX-W-WDGWIS	
BNL2 FD2	р <u>+</u>	, œ	LSI	
HC-J6	2a	231		
HC-J8	2p	232	S-NTWOLTVL	
CH610	3c	233	WQLEG-V	
S83	3c	254	1 1 1 1	
NE92	2d	234	QWQLRVVEEKI	
BNL3	2e	16	WQLXVVENSSGRFH	
FR4	2£	18	-MAA-DWQLRVVE-SRTFT-VS-NVSR	
BNL4	2g	20	-MAS-NIWQMQG-VVELQKIPVNVNQ	
BNL5	5h	24		
BNL6	2i	56	EWKD-T	
FR13	2ķ	9/	-MS-XTWQLXX-VVESIPVX-XVSR	
BR36	3а	255	-VLSD-VIQDT-TTPVVKY	
HCV-TR	3p	235	-VLS-GE-VLTTQ-STTVSTV-T	
24	4a	256		
GB809-4	4a	257	ITDHHLAVTPVAVS	
21	4p	258	TTEHH-M-LTE-TPLAPY	
GB116	4°C	241	ILAPY	
GB215	4c	2	LSAPY	
GB358	4°C	260	IL	
DK13	4 d	236		
CAM600	4e	237	IATENHLTQLSPY	
GB809-2	4e	238	IATDNHLKTQLSPY	
CAMG22	4 £	261	L	
CAMG27	4 £	262	II-LEHHLT-QI-LL-APH	
GB549	49	243		
GB438	4 h	244	IPLVPY	
BNL7	4 k	30	-YLAPY	
BNL8	4 k	32		
BNL9	4 k	34	IDHHLVQ-SLI-APY	
BNL10	4 k	36		
BNL11	4 k	38		



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BNL12	4]	40	LAPI
GB724	4×	246	IAVZ
BE95	5a	5a 239	
BE100	5 a	263	
HK2	6а	240	LLDAMLLVDDR-TH-VL-IPN
VN4	7c	44	LQASL-VPN
VN12	7d	48	LNGMLKTLTKLSASL-VQN
FR1	9a	42	LS-NFETMLIKAELPVSL-VPN
NE98	10a	52	-MS-GTSTTXx



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4
Fig

Isolate	Туре	SEQ ID	V5 V5
HCV-1	1a	229	PATQLRRHIDLLVGSATLCSALYVGDLCGSVFLVGQLFTFS
HCV-J	9 7	230	A-A
BNL2	1 g	r 00	T-AFRM
FR2	1£	12	A-VFM-IGT
HC-J6	2a	231	G-M-AA-M
HC-J8	Sp	232	-AV
CH610	3c	233	
S83	3c	254	PGALTKGAII-MVVALM-AA-VVVVQH-TF
NE92	2d	234	I
BNL3	2e	16	^ ^
FR4	2£	18	PGALTRGATI-MIA-MIAA-VAVVQY-TF
BNL4	2g	20	PGALTRGTTI-MVIVA-MIAA-VVIVQH-NF
BNL5	5h	24	
BNL6	2 i	56	PGAXTKGTII-AF
FR13	2k	9/	PGALTEGSTI-AFIVAIMIAA-VVIVEH-HF
BR36	3а	255	VGATTASI-S-VAMMAAQ-
HCV-TR	3b	235	LGVTTASI-T-V-MARQAF-AAT-
24	4 a	256	PGA-LESFVMA
GB809-4	4a	257	MDA-LESFVMAVVGAMQ
Z 1	4p	258	PNA-LESMVMAMF-IGD-R
GB116	4c	241	VGA-LESS-VMAVIGM-S-Q
GB215	4c	259	IGA-VESFV-MMAVIGM-S-R
GB358	4c	260	IGA-LESS-VMAAIGM-S-Q
DK13	4 d	236	LNA-LESVMGIVGQ
CAM600	4 e	237	AGA-LEPVMAMIGLMQ
GB809-2	4 e	238	VGA-LEPVMAVGLMQ
CAMG22	4 £	261	LGA-LESMVMTGIAMRL
CAMG27	4 £	262	IGA-LESMVMTIGIM-N-RL
GB549	49	243	VGA-LESMVMAVIGMR
GB438	4h	244	LGA-L-SV-Q-VMAIHGAMVS-Q
BNL7	4 k	30	VIX-XGL
BNL8	4 k	32	WN-S-
BNL9	4 k	34	-S-VMAVIGA
BNL10	4 k	36	TAA-LESS-VMAVI-XGLM-SXQ
BNL11	4 k	38	IGA-LESS-V-VMAVIGLM-S-R



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LSA-LMSVVM						ASVSIRGV-E-VA-AFMGLRMYEI		
40	246	5a 239	263	240	44	48	42	52
41	4×	5a	5a	6a	7c	7d	9a	10a
BNL12	GB724	BE95	BE100	HK2	VN4	VN12	FR1	NE98



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V5 205	TOGCNCSIYPGHITGHRMA	V-B V-DXXX V-DT	V-EX	- D	V-EX S-D	1 1	V-TVS	DT	DAV	DAV	DA		- 1	DV	A-D	D		V-D
SEQ	229 230 4	8 12 231	232 233 254	234 16	18 20 40	76 255	235 256	257 258	241 259	260	237	261	243	244 30	32	34 36	38	4 ○
Туре	la ld	1d 1f 2a	20 20 20 20	5 g g	2g 2g	3a 3a	3b 4a	4a 4b	24 0 4	4°C	4 4 e	4£	. 4 .	4 4 다 차	4 k	4 4 X X	¥.	4 T
Isolate	HCV-1 HCV-J BNL1	BNL2 FR2 HC-J6	HC-J8 CH610 S83	NE92 BNL3	FR4 BNL4 BNI,5	FR13 BR36	HCV-TR Z4	GB809-4 Z1	GB116 GB215	GB358 DK13	CAM600 GB809	CAMG22	ו ס	GB438 BNL7	BNL8	BNL9 BNL10	BNL11	BNL12



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V-N-----S--V--V-D-----S--V--V-D-----T--V--V-E-----T----A-D-----A----246 239 263 240 448 42 52 4x 5a 5a 6a 7c 7d 9a GB724 BE95 BE100 HK2 VN4 VN12 FR1 NE98

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56/74

GB2154c273-----T--A--C--A-AA-----A-GGTC-----AGG-A--T--G GB3584c274-----T--A--C--A-AG-----A-GGTC-----AGG-G-T--G-GB8094e275-----T--G-----A----AAGGTC--A--A-A-G----T--G-GB54949276-----G--G--C--A--G--T---A-G--C-----A-AG-------G-BNL84k83-----T--A--C--A-AG-----A-GC-C----A-AGG---T--G-BNL124185-----G--G--C--A-AG-----A-GGTC----A-AG----T---NE9810a97----T----TGG------A-GGTA--ACTTT-C---TT--GG EG814m87-----C--A--C--A--G-----A-GGTC-----AGG----T--G-FR19a95A------G-G-C-----A--C--ACNA-AC-T---TG-CHR185a277---G--C--T--C--ACAT-----AATG--T--A---T-T--T-----A----G----A--G--A--A--AT-C GB1164c272----T--A--C FR1911a105---T--T FR1411a99---T--C FR1511a101---GB484c271---7932 7981

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57/74

HCV-11a264GTTGTGACCTCGACCCCCAAGCCCGCGTGGCCATCAAGTCCCTCACCGAG HCV-J1b265-----T-G-C---G----A-GCA-----A-G---A--PAK643q81------T--G--GG-G--TA-ACG---A----A--G----A CAM10781e61-C-----GC----G----A--T--A-----TT-G--T FR21f63-C----T--A-----G-G--T---AA-----A-G------A-G-----T---FR161g67-C------G-C----G-G-T----A-----A-----T----HC-J62a267C----TC-T-GCC-GAGG-G----A-ACT----AC-C--A-G--T-HC-J82b268C----TCT--GCCT-AAG-----A-AACT-T---AC-C--G----T-BE901b266-----T-G-C---G-G---A-ACA-----A----A-BNL52h73CC--CTC-T-ACC-GAG-----AACT-----AC-T--AT-G--T--FR132k77CC---TCA--TCC-GAGG-G----A-CT----AC-C--A--A--T--FR182179CC--CTCGT-GCC-GAGG-G----GACT-T---AC-T--G----T-BNL32e69C----TC-T-ACC-GAG--G---A-AACT-----AC-C--AT-G--T--FR42f71CC--CTC-T-ACC-GAG--G-----GACT-----AC-T-AT-A--T--BNL111d53------G-----BNL21d55-----AA-----AA-----G------BNL21d55------T-G-C---YG-G-----AA-----AA------G------FR171d57-C-----T-G-C----G-G------AA------AA------G------T13a269-C---A---T--A--GG-G---A-GAGA-TG---TCC------G-79828031

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CHR185a277CA-TGT--T-GC-G--TG-G--T-----A--ACG---A---C-A ---G--G---G------AA--TA--T-CCG----A--A--VN137a89-C---A-GT-G--G--GC---A--GACA------CA--G--T--T--C NE9810a97CC-----A-GGA-G-G--TA-GAG--TG--A-CT--A-----G--FR1511a101C----C-AT-GCC-GAAG-G-----GAA-----T--A--A--A-BNL124185-----G--R--G--R---G------AAA--A--ATCCG------A--EG814m87-----T-G--G--AG-G-----AA---A-ATCCG-----G--FR1911a103C----C-AT-GCC-GAAG-G-----GAA-----A--A--G--A-VN127d93-C--CC-AT-A--T--GGT---A--GAAA-----T-CA--T--T-T-VN47c91-C--CC-A--T-----GGTG--A---AA---T--T-CA--T--G--T--FR1411a99C----C-AT-GCCTGAAG-G-----GAAA----T--A--G--G--A-GB2154c273-----GB484c271-----79828031



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HCV-11a264AGGCTTTATGTGGGGCCCTCTTACCAATTCAAGGGGGGAAAACTGCGG HCV-J1b265C------C----T--C--G--T----G-A----C-----BE901b266C-----A-C----T--C--G--T-----A----C----T--BNL52h73--A----C--A--A--G--CA-G-TG--CAGC-A---AC--TC----FR132k77----G--C--G--A--T--CA-GCAG--CAGC-A----C-ATC-----FR182179--A--C--CA-A--C--G--GA-G-TG--CAGC-AA--CC--TC---T T13a269C-----CTGC-----A-GTT---CAGC-A-----CCC-A--T-T93b270C----G--CA-C-A--T--CA-GTA---CAGT-A----CTCC-G---BNL32e69--A--C--C--A--A--G--CA-G-TG--CAGC-AA---C-ATC-------C----A--T--CA-GTT---CAGC-A----CTC--A--FR42f71--A----C--G--A--G--CA-G-TG--CAGC-AA---C--TC----CAM10781e61----G-C--C----G--CT-G-------AA-----FR21f63-----A-C----T--C--G--A--C---AA---C PAK643g81C---80328081

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60/74

CHR185a277C-C--G--CTG---A----CA-GTAT--CAGC-A----C-AC-A--T VN137a89C-AT-G--CTNC--T--T--CA-GTNT--C--T-AA--TC--GCA--T--BNL124185--R--C--CT-G--C----CA-GTAT--CAGC-AA------CT----EG814m87--A--C----G--C-T--CA-GTTT--CAGC-A---A--CCTA--T-VN127d93C----G--CTGC--C----CA-GTA---C--TC-A--TC--TCA--T FR19a95-----C-----C-----A-GTA---C---A---CC-ACT---T FR1511a101--A--A--C--G--C---GA-GGAA--CAGC-AA--CC--GC--FR1911a105--A--A--C--G--C-----GA-GGAA--CAGC-A---CC--GC--VN47c91C----G--CTGC--W--G--CA-G-TG--C--CC-T--TC-ATCA--T

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=ig.5G

HCV-11a264CTATCGCAGGTGCCGCGCGAGCGGCGTACTGACAACTAGCTGTGGTAACA BE901b266-----C-A-----A----G----G-----G-----T HCV-J1b265T-----C-----A--T----G-----G-----C--C--C FR161g67------C-----T-----T--G-----T--C------HC-J62a267G--CA-GC-T--C----ATG-G-HC-J82b268---CA-GC-T--C----ATG-G-HC-J82b268---CA-GC-T------A-G--G--T--C--C--A-TG--G-BNL21d55-----TC----T----T-----G-----C--C----BNL52h73T--CA-AC-T------C----A--G--C--C--C--ATG--G--T CAM10781e61------A----T--C-----CT-----C BNL32e69A--CA-GCAT----BNL11d53---C--TC-----F93b270----C-C-C-**F13a269T----**C PAK643g81A---

80828131

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62/74

----A------CTAC--C--C--TC--G--NE9810a97T--C---C-C----T--T--G--G--AC-C--C----TC--G--FR1411a99A---A-GC-T------A-G--T-C--C--A----TG--G-----A-----G--T-C--C--A----TG--G-----C-----CT-C--C----TATG--C VN137a89A--C--T----A-G--C--T-----CT----C-C-T-CTG-CC--VN127d93G--C------G--T--T--CT-C--C--A----TG--C-FR19a95TC-A---C-A----A-A----A-----CC-C--A---ATG----GB484c271G-----A--T----A-----GB1164c272G-----A----80828131 GB8094e275G----T--A----GB5494g276GC-A--G-----BNL84k83G----G--A----BNL124185G----G------FR1511a101A---A-GC-T----FR1911a105A--CA-GC-T----CHR185a277T----T--A--GB3584c274G----GB2154c273G----



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FR161g67-----T---A-----C-G--A---GCC--G-------AAG----HC-J62a267--A----TG-G--A---TTA--G-----AAG--T-----A-A HC-J82b268--A-G--A--T-----A---TT-----G---AAG--T----A--HCV-11a264CCCTCACTTGCTACATCAAGGCCCGGGCAGCCTGTCGAGCCGCAGGGCTC HCV-J1b265-----T--A--T---T-G-----ACT--G-------T---AA---BE901b266----T--A--T---C-A-----TCT---------T--GAA---BNL21d55-----T-G-A----T-G-A--A---G-------T-GAA----FR171d57------T-G-A--T-G-A--A--------T-GAA----CAM10781e61------C----T-----TA------A----T--CAA-FR132k77-T----A-----T-G-----A---G-----CA-G-----G--CA-T FR182179--A---G----TG-G--A---AT-----T--CA----T--C--A-T BNL11d53----T-----T-G--A---A-A--G-------T---AA---FR21f63-----T----A----A-----T--GAA--------ACT----A-CA-G--T--G--T--81328181

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FR1511a101--A-G----T------AAR-----T---AA---Y--T--CA-T NE9810a97-AA----C--T------A--AAA-----TACCAA---T--C--AA-T FR1411a99--A-G----T-----A--TAAA--G--T---AA----T--CA-T VN127d93--A-G--A----C-G-----T-A---G--T---A-G--A--RAA---FR19a95--A----A----T-C-G-----AACC--C--T--C----A--C--CT-T GB484c271-A--G--G-----C----A---TCA--C--TATCAA---G--G---GB1164c272-A--G--G----TC----A---TCA--C--TATCA----G--G-GB2154c273-A--G--G-----TC----A---TCA--C---ATCA-G--GT--81328181 FR1911a105--A-G----

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HCV-11a264CAGGACTGCACCATGCTCGTGTGTGGCGACGACTTAGTCGTTATCTGTGA PAK643g81--A---CCAT-AT-C--T--C--C--A--T--T--G--G--AG-GGC-T13a269-G-A--CCGGA-T-T--T--C--C-A--T--TC-G----AG-GGC--T93b270A-A---CCAT-TT-C--T--C--C-A----T--G--G--G-A-C--HCV-J1b265------G-----AAC--A-----C-T---------C--A----C-T-------C--A----C-T----BNL21d55---------FR171d57-------A---CAM10781e61-----



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=ig.5L

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Fig.5M

82328271 HCV-11a264AAGCGCGGGGGTCCAGGAGGACGCGGCGAGCCTGAGAGCC HCV-J1b265G--T----AAC-----T-----GC--AC----BE901b266-----AAC---A---------AC---T BNL11d53G--T----A---G----A-------AC----T BNL21d55G-----A---G--------A---AC---T FR171d57G--T--R--A---G-----T-----A---AC---T T93b270----TGC--C---G-----AGA--AGCT---C--PAK643g81G--TTGC-KC--TG-T----G-ATAG-GCAGC CAM10781e61G--T-TA---AC-----T----T13a269G----AT--C---G-T-----TAGA--AGC---BNL52h73G--TCA---AAC-G------T-A-CG--A---FR132k77G--TCA----ACTG--AG-----A-AAC-A---FR182179G--TCA----AC-G-----A-CGA-AT--FR161g67G--T--------T--T--A---HC-J62a267G---CA----AC-G-------A-CG--A HC-J82b268G---CAA--TAA-G-------A-CGA-A BNL32e69G--TCA---A---G------ACCG--A-FR42£71G--TCA-----CTG------A-CGA-A--FR21f63G-T--A---N----TC--T---



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Angle with the

2645	STVTESDIRTEEAIYQCCDLDPQARVAIKSLTERLYVGGPLTNSRGENCG	0K	NK-Q; NVSA-EKIK-Q;	A	XXV-XSA-EK-Q	RSRA-S-PEE-HTHMFK-QT	RK-QSA-S-PQETV-HMK-QS	S-S-PEETHK-QS	RSLA-S-PETHMLK-QT	RXSA-S-PETHMMK-QS	RSLA-S-PETHMMK-QS	A-RSLA-S-PETHMMK-QS	RVSV-LS-S-PEEAHMQK-QS	RNS-FLA-S-PEETV-HIMMK-QS	MFK-AQ	K-AQ	K-AQ	HE-E-E-E-E-KSAIMYK-LQ	QVEE-E-ER
SEQ ID	278	280 54	28 28	62 64	89	281	282	283	284	70	72	74	78	80	285	286	287	288	82
Type	la 1b	10 10	1d	le 1f	19	2a	5 p	3c	2d	3e	2£	5h	2 7 2	21	3a	3а	3a	3p	39
Isolate	HCV-1 HCV-J	2TY4 BNL1	BNL2 FR17	CAM1078 FR2	FR16	HC-J6	HC-J8	ARG8	NE92	BNL3	FR4	BNL5	FR13	FR18	BR34	BR36	BR33	T9	PAK64



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	DIWH	WHK-DL	MHK-DL	WHK-DL	DI	MYK-DL	KMYK-DL	WHK-DL	WFK-DL	WH K-DI	WHK-DL	LMYKL	WFK-DL	CMXK-00	CMYK-00	XMXK-OA	CXMM0S	CMYOS	MYK-OL	CMFK-OH	MEK-OA	MEK-QA	10 22
9KVEVE-EKTAMHK-DL	E-ERTA	E-EKV-TA	E-EKTA	E-EKV-AA	E-ET-KV-SA	E-EKV-SA	E-EKV-SA	N-EXDX-KV-NA	E-EKV-TA	N-E-EKTA	E-EKV-TA	X-EKSAX	E-E-KSA	O-EARO	.Y-Q-ERQ	K-E-ATTD	QVKT	QVKT	OEK	KDERV-T	Q-PEEK	Q-PEEK	7 555
KVEV	KVEV	KV	KVEV	RKVEV	RVEV		RVE	PR-X-VEV	X-RGEV	A	KPEV	KVE	RVEV	-SSMH	ISSWH	R-VQHD	RXHD	-SRHD	GRXDLS-	QV-LS-F-A-	RB-	RS-XXA-	4 1 20 C
289	290	291	292	293	294	295	296	297	298	299	84	98	88	300	301	90	92	94	96	98	100	102	301
4°C	4c	4°C	4°C	4e	4 £	49	4ĥ	054i	1 4j	4.2	4 k	41	4m	5 a	5a	7a	7c	7d	9a	10a	11a	11a	110
GB48	GB116 4c	GB215	GB358	GB809	CAMG22	GB549	GB438	CAR4/12	CAR1/50	EG13	BNL8	BNL12	EG81	BE95	CHR18	VN13	VN4	VN12	FR1	NE98	FR14	FR15	0100

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	respective to the second of th	1 1		· · · · · · · · · · · · · · · · · · ·			MIVLKIIAPS-	FMKLKIV-PVS-	MVV-	MIVQKIIAPS-	1	K	MIVLKIVAPI-S-	MLQIVAPS-	-FMIVMIDAPS-	?FITARNPDFVA-	2FITAKRSPDFVA-	9FITAKRNPDFVA-	PFIKSK-PSF:VS-
2605	YRRCRASGVL				1 1 1 1 1					E			1 1 1 1 1 1 1 1 1	1 1 1 1 1 1 1	(±)	1	1		1 - : :
SEQ ID	278	280	200	62 62	64	89	281	282	283	284	70	72	74	78	80	285	286	287	288
Type	1a 1b	15	, 1d	je Je	1 £	1g	2a	3p	3c	2d	3e	2£	2h	2k	21	3а	3а	3а	3b
Isolate	HCV-1	2TY4	BNL2	FK1 / CAM1078	FR2	FR16	HC-J6	HC-J8	ARG8	NE92	BNL3	FR4	BNLS	FR13	FR18	BR34	BR36	BR33	T9

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GB48	4c	289	1 1 1 1 1	X	E	T	S	-IK	RA-
GB116	4c	290		Y1		I	-S-	I-	.RA-
GB215	4c	291	1 1 1 1 1 1	Y1	· ·	T	-S	SI	.RA-
GB358	4c	292		Y1		I	S-	I	.RA-
GB809	4 e	293	1 1 1 1 1	Y1	W i	T	S	I	.KA-
CAMG22	4 f	294		Y1	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	FL-	Ė	-TK	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
GB549	49	295	ō	Y1	YKG	1	- ^-	L-	S-
GB438	4h	296	L	Y1	^\	I	Ŀ	I -	.KA-
CAR4/1205	.4 i	297	II	Y1		1	Ė	T	. K A -
CAR1/501	4 j	298	0	F1		I	L-	L	- K
EG13	4.2	299		F1	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	I	.T.	I.	-R
BNL8	4 k	84	1 1 1 1 1 1 1	Y1		1	SS	I	RA-
	41	98		Y1	^	I	Į.	. T	- K A -
	4m	88	1 1 1 1 1 1 1	Y1	بغ	1	Į.	L	K
	5a	300		FF	4M	!	-I-S	3R-	RLA
	5a	301		JB	4M		S-1-	-L-SK	BB
	7a	90	1 - 1	II	Ar	1	ģ		K-FDS-
	7c	92				T	ģ	K-	K-KNYDA-
	1d	94		F1	W	I	ģ	XK-	KNFDA-
	9a	96	00	1d-	I	FL	Į.	4	FT-YDVT-
	10a	98	1 1 1 1 1 1	- b I	I e	1	- X	TKI	KNPSFA-
	11a	100	1 1 1 1 1	-EI	W	1 1 1	. .	KIV-PV-	V-PVS-
	11a	102		F1	W		X-	KX I	IV-PVS-
FR19	11a	106	1 1 1 1 1 1 1	F1	W	1	X	XI	V-PVS-

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Fig. 6E

	2745 2757	SAGVQEDAASLRA	T	TV	N	EV	-XEV	-V-T	IE-XXPS	1 1 1 1 1 1 1 1 1 1	-Q-TEERN	-Q-NEERN	-TE	-QEDRN-	-Q-AEERNV	-Q-TEERNV	-Q-TER-ENNP	-Q-TEERNV		1	1	-CER-A	-CX-D-EDRAALR
SEQ ID		278	279	302	54	99	28	62	64	89	281	282	284	70	72	74	78	80	285	286	287	288	82
Type		1a	1b	1p	1q	1q	1d	1e	1 £	19	2a	5 p	2d	2 е	2£	2h	2k	21	3а	3а	3a	3b	39
Isolate		HCV-1	HCV-J	BE90	BNL1	BNL2	FR17	CAM1078	FR2	FR16	HC-J6	HC-J8	NE92	BNL3	FR4	BNL5	FR13	FR18	BR34	BR36	BR33	T9	PAK64



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GB48 4C 289 -DEKRP-G-GB116 4C 290 -DEKRP-G-GB215 4C 291 -DEKRA-G-GB358 4C 292 -DEKRA-G-GB89 4e 293 -DEKRA-G-GB89 4e 293 -DEKRA-G-GB49 4g 295 -G-EKRA-G-GB43 4h 296 -G-ERA-G-GB43 4h 296 -G-ERA-G-GB1 4i 297 -I-IDKQA-T-CARI/501 4j 298 -D-ERA-G-BNL2 41 86 -EEKRA-Q-BE95 5a 301 -O-THE																							
4C 28 4C 28 4C 29 4C 29 4C 29 4B 29 4D	EKRP-	EKRA-	EKRA-	-EKRA-	EKRX-	DERRA-	GERA-	GERA-	I-IDKQA-	EPXTX	DENRA-	EESQP-	DRRA-	Q-THE	-THK	LSTSA-	GSVDA-	PGA-	N-IN	-ID	KE	KEQRD	EORD-
115 115 115 115 115 115 115 115	œ	9	σ	6	σ	σ	σ	9	σ	σ	4	ø	œ	0	0	0	95	94	96	86	Õ		ō
	4°C	4°C	4 _C	4c	4 e	4 £	49	4h	054	1.4	4 ,	41	4m	5a	5 a	7a	7c	7d	9a	O	-	-	-
	GB48	GB116	GB215	GB358	GB809	CAMG22	GB549	GB438	$\overline{}$	in	m	BNL12	EG81	BE95	CHR18	VN13	VN4	VN12	FR1	NE98	FR14	FR15	FR19